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Specialist Field¹: Statistical Learning for Big Data; Bayesian Computation and

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Instructions

I have been instructed to provide my opinion in respect of the following issues:-

- 1. The appropriateness of using binomial analysis as suggested by MTRCL in investigating the defective rate of coupler connections at Hung Hom Station Extension ("HUH").
- 2. The rationale and considerations in relation to the random sampling of coupler connections at East West Line ("EWL") and North South Line ("NSL") slabs for the Stage 2 opening-up investigation under the Holistic Proposal.
- 3. The statistical analysis of the results of the enhanced Phased Array Ultrasonic Test ("PAUT") obtained from the opening-up investigation for deriving the coupler defective rates for EWL and NSL slabs.
- 4. The statistical analysis to derive a combined defective rate to account for the condition on both sides of coupler connections at those locations where the EWL slab was connected to the diaphragm wall ("D-wall") via capping beam.

¹ A copy of my CV is attached as Appendix A.

Declaration

I declare that:-

- (a)I have read the code of conduct set out in Appendix D of the Rules of the High Court (Cap. 4A) and agree to be bound by it;
- (b)I understand my duty to the Commission; and
- (c)I have complied with and will continue to comply with that duty.

Professor YIN Guosheng

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Date: 16 September 2019

Opinion

1. Appropriateness of using binomial analysis as suggested by MTRCL in investigating the defective rate of coupler connections at HUH

1.1 Background

- 1.1.1 The Shatin to Central Link ("SCL") is a railway project of Hong Kong under construction, which involves construction of new stations and the extension of existing stations (including the HUH). The Government entrusts to the MTR Corporation Limited ("MTRCL") the design, construction, procurement of services and equipment, testing, commissioning and all other matters associated with the bringing into service of the SCL. The construction work for HUH is being carried out under Contract No. 1112. The Contractor of Contract No. 1112 is Leighton Contractors (Asia) Limited ("Leighton").
- 1.1.2 At the HUH, two new platforms were constructed in the form of two reinforced concrete slabs supported by D-walls. The reinforcement connections between the platform slabs and the D-walls were formed by screwing rebars of the platform slabs into couplers embedded in the D-walls.
- 1.1.3 In May 2018, there were media reports that there had been unauthorized cutting of threaded ends of rebars during the construction of the above-mentioned platform slabs. Such allegation, if true, casts doubt on the quality of workmanship on whether the rebars are properly connected to the couplers embedded in the D-walls.
- 1.1.4 MTRCL had then been following up by reviewing the available photographic and other site records of the construction works of the platform slabs. MTRCL appointed independent expert consultants for a holistic study which was intended to verify the as-constructed condition of the HUH of the SCL Project.
- 1.1.5 After several rounds of discussion between the Government and MTRCL, MTRCL submitted to the Government on 4 December 2018 a Holistic Proposal for Verification & Assurance of As-constructed Conditions and Workmanship Quality of the HUH (EWL Platform Slab, NSL Platform Slab and the Connecting Diaphragm Walls) ("Holistic

- Proposal"). The Holistic Proposal is a staged approach exercise consisting of Stage 1 (desktop exercise), Stage 2 (physical investigation) and Stage 3 (structural assessment).
- 1.1.6 In particular, the Holistic Proposal includes sampling method for verification of workmanship of coupler connections. Considering the specialized nature of the sampling design and the need to ensure that findings would be statistically meaningful, I and my two colleagues of the Department of Statistical and Actuarial Science HKU ("HKU Statistics Team") were engaged by the Government to provide independent statistical advice and expert opinions on MTRCL's Holistic Proposal in respect of its adequacy and any deficiencies in the sampling approach, methodology and associated statistical analysis throughout the process of sampling and subsequent analysis.

1.2 Proposal to use binomial analysis

- 1.2.1 The proposal to use binomial analysis was initiated by MTRCL in November 2018 in a draft Holistic Proposal for the purpose of assessing the workmanship in the coupler connections and rebar fixing in light of the allegations raised. It was suggested the extent of opening up should be based on statistical theory. A sample size of not less than 84 randomly selected coupler connections will give a meaningful result with 95% confidence level using binomial statistics. The coupler/rebar connection process was assumed to be similar at the EWL slab and NSL slab and the occurrence of defective coupler connections was considered as random in general. Therefore, the test results of coupler connections at one location can be treated as independent representative results for statistical analysis. I was given to understand that the construction method and design details of the EWL and NSL slabs are different. The EWL and NSL slabs are therefore treated as two populations.
- 1.2.2 I was asked by the Government to comment on the methodology proposed by MTRCL and to verify the accuracy of the sample size calculation provided.
- 1.3 <u>The underlying assumptions for the binomial distribution and</u> appropriateness of binomial analysis

- 1.3.1 In the first phase, I considered whether binomial analysis was an appropriate method by going through whether the underlying assumptions were satisfied.
- 1.3.2 As its name would suggest, a binomial distribution means each trial will lead to two possible outcomes, say either pass or fail. Suppose a random sample size of n coupler connections is drawn for assessment. Each selected coupler connection can have two possible outcomes only, either pass or fail in terms of the quality of the workmanship, and the coupler connection of a failed quality of workmanship is considered defective. The whole exercise focuses in the estimation of p, i.e. the proportion of defective coupler connections in the population.
- 1.3.3 The binomial distribution is frequently used to model the number of "failures" or "successes" in a random sample of size *n* in clinical trials and statistical quality control. The binomial analysis is only appropriate if the following assumptions are satisfied:-
 - (i) The experiment consists of n identical trials;
 - (ii) Each trial results in one of the two possible outcomes, either "Defective" or "Non-defective";
 - (iii) The probability of selecting a defective coupler connection in each trial equals to p and remains the same;
 - (iv) The trials are independent.
- 1.3.4 The random variable of interest is the number of defective coupler connections observed in the *n* trials. The appropriateness of the four assumptions listed above was discussed in consultation with the Government's project team.
- 1.3.5 First, assumption (i) is satisfied naturally as the opening-up exercise to expose the coupler connections for each of the *n* selected locations is identical, regardless of the location of the subject coupler connection. Second, as we only considered the quality of the workmanship on each coupler connection to be either satisfactory (i.e. complying with the supplier specification) or not satisfactory (i.e. failing to comply with the supplier specification), assumption (ii) is also satisfied by the nature of the opening up exercise. Furthermore, the coupler connections in question were carried out by the same contractor, under similar site condition and the quality of workmanship is considered to be generally consistent with respect to the locations of the coupler connections. In

other words, the defective coupler connections are random events which are distributed randomly, and the probability of selecting a defective coupler connection can be regarded as constant. Moreover, the outcome of a coupler connection will have no effect on the outcomes of other coupler connections; hence assumptions (iii) and (iv) are also satisfied. As such, binomial analysis is considered a reasonable and suitable approach for the purpose.

1.4 Adequacy of sample size

1.4.1 With the assumptions satisfied, I now discuss the adequacy of the sample size suggested by MTRCL. A general practice is to use a confidence level of 95% throughout the sample size calculation exercise. I calculated, p_U , the upper bound of the 95% (1-sided) confidence interval for the proportion of defective coupler connections in the population using the exact binomial probability formula:

$$\sum_{k=0}^{y} {n \choose k} [p_U]^k [1 - p_U]^{n-k} \le (1 - 0.95)$$

where n is the sample size, y is the observed number of defective coupler connections in the sample. A higher accuracy is associated with a smaller margin of error $(|p_U - p|)$ leading to a larger sample size required. In other words, a smaller sample size will yield a larger margin of error. For example, the estimated margin of error is estimated to be 5.8% when n = 50 and y = 0.

- 1.4.2 As suggested by MTRCL, a random sample of size n = 84 yields an estimated margin of error of 3.5% when the number of defective coupler connections in the sample is zero (y = 0). In other words, if none of the 84 exposed coupler connections are found to be improperly connected, no more than approximately 3.5% of coupler connections in the population could potentially be defective (in a worst case scenario with 95% confidence level). This estimated margin of error (i.e. the maximum failure rate) is considered to be reasonable and acceptable, with due consideration of cost and time implications.
- 1.4.3 The table below shows the maximum failure rate in the population based on the binomial statistical approach for a total number of samples of 84.

Total sample number = 84	
Total number of failures in	Maximum failure rate in the population at
the samples	95% confidence level
0	3.5%
1	5.5%
2	7.3%
3	9.0%
4	10.6%
5	12.1%
10	19.4%
20	32.7%
30	45.2%

2. Rationale and considerations in relation to the random sampling of coupler connections

- 2.1 With the submission of the final Holistic Proposal by MTRCL on 4 December 2018, I was invited to conduct the random sampling exercise for the locations to be opened up. This section is to illustrate the methodology for the selection of coupler samples for the Purpose (ii) investigation, which is to verify the workmanship quality of the coupler connections between the D-wall panels and EWL/NSL slabs in Areas A, HKC, B, C1, C2 and C3. The methodology was designed and the random selection was conducted by the HKU Statistics Team led by myself.
- 2.2 <u>D-wall panels available for selecting sampling units at EWL and NSL slabs</u>

EWL slab

2.2.1 The EWL slab is connected to East D-wall and West D-wall of approximately 400 metres run from Gridlines 0 to 50, comprising a total of 234 D-wall panels. These D-wall panels to EWL slab connections can be divided into four groups, namely EWL East Wall Top connections, EWL East Wall Soffit connections, EWL West Wall Top connections and EWL West Wall Soffit connections. Rebars connecting D-wall panels and EWL slab in these four groups of connections include design with coupler connections or suspected to have coupler connections or with

straight continuing rebars only.

2.2.2 Before conducting random selection of coupler connection samples for verification, the Government and MTRCL, after going through the relevant construction records, reached general consensus on the identification of D-wall panels with or suspected to have coupler connections among the 4 groups of connections. The number of D-wall panels identified to have coupler connections are summarized below:-

Group of connections at EWL slab	No. of D-wall panels with / suspected to have coupler connections
(A1) EWL East D-wall Top connection	27
(A2) EWL West D-wall Top connection	10
(A3) EWL East D-wall Soffit connection	88
(A4) EWL West D-wall Soffit connection	107
Total :	232

NSL slab

- 2.2.3 Similar to EWL slab, the NSL slab is connected to East D-wall and West D-wall but at a level lower than the EWL slab. The D-wall panels to NSL slab connections can also be divided into four groups, namely NSL East Wall Top connections, NSL East Wall Bottom connections, NSL West Wall Top connections and NSL West Wall Bottom connections. According to the details in the original working drawings (which as I understand have been complied with), rebars connecting D-wall panels and NSL slab are all connected by couplers.
- 2.2.4 The Government and MTRCL had gone through the relevant construction records before the random selection exercise of coupler samples at NSL slab for verification. MTRCL advised that certain locations of D-wall panels and NSL slab connections are physically inaccessible for verification of coupler connections based on the following reasons:
 - (i) the NSL slab is founded on soil which made the slab-to-D-wall bottom connections at NSL slab physically inaccessible from the underside of the NSL slab; and

- (ii) the presence of mass concrete filling on top of the NSL slab has completely filled and covered up NSL D-wall Top connections at certain locations in Area A and Area B, which also made the slab-to-D-wall top connections at these panels physically inaccessible.
- 2.2.5 In view of the constraints mentioned in paragraph 2.2.4 above and as jointly agreed by the Government and MTRCL, the D-wall panels at which coupler connections are available for random sampling for Purpose (ii) investigation and the amount of these D-wall panels are summarized below:-

Group of connections at NSL slab	No. of D-wall panels with / suspected to have coupler connections
(B) NSL East and West D-wall Top connection	189

2.3 Methodology of two-phase cluster sampling scheme

- 2.3.1 The aim of the random selection of coupler connection samples (conducted independently by the HKU Statistics Team) was to select from the overall population of coupler connections in the D-wall to EWL and NSL slabs connections the locations at which the coupler connection samples are to be opened up for verification of workmanship.
- 2.3.2 A two-phase cluster sampling scheme was adopted in the selection of sampling units, each opening-up site (or sampling unit) yielding three coupler connections. Phase 1 sampling selection was to determine the locations of sampling units on plan, while Phase 2 sampling selection was to determine the layer of coupler connections to be exposed for workmanship verification at locations selected in Phase 1. The methodology of the sampling selection is discussed in detail in the following paragraphs.

Phase 1 sampling selection

2.3.3 As described above, the Government and MTRCL jointly identified 232 and 189 D-wall panel locations at EWL slab and NSL slab respectively, which are physically accessible for site verification, and thus available for Phase 1 sampling selection. Based on the prior decision made, 28

sampling units, each yielding 3 coupler connections, would be selected from each of EWL slab and NSL slab.

2.3.4 For EWL slab, the top connections available for sampling were significantly fewer than those at the soffits. It was considered more appropriate to select sampling units at each group of connections separately on a proportional basis to ensure the sampling units selected would be more proportionally distributed in the 4 groups of connections and that random samples from all 4 groups will be selected (to enhance representability of the samples). The number of sampling units to be selected from D-wall panels in each group of connections are tabulated below:-

Group of connections	No. of D-wall panels with / suspected to have coupler connections	No. of sampling units selected
(A1) EWL East D-wall Top connection	27	3
(A2) EWL West D-wall Top connection	10	1
(A3) EWL East D-wall Soffit connection	88	11
(A4) EWL West D-wall Soffit connection	107	13
Total	232	28

2.3.5 For NSL slab, the numbers of top connections available for sampling at East D-wall (92 panels) and West D-wall (97 panels) were roughly the same. Therefore, it was considered unnecessary to select sampling units from these two groups of connections on a proportional basis. These panels were pooled together for random selection of sampling units for NSL slab.

Group of connections	No. of D-wall panels with / suspected to have coupler connections	No. of sampling units selected
(B) NSL East and West D-wall Top connection	189	28
Total	189	28

- 2.3.6 In order to select D-wall panels on a random basis, a number with 5 decimal places was randomly generated from a uniform distribution ranging from 0 to 1 and assigned to each D-wall panel in the group. D-wall panels available for selection in each group were then sorted in a descending order based on the assigned random number, and the required number of D-wall panels were selected from the top of the list, i.e. with the largest random number, downwards. The D-wall panels listed after the required number of selected D-wall panels formed the "waiting list" and served as back-up replacement locations in case difficulties were encountered during opening up of the coupler connections at the originally selected D-wall panels. For instance, for EWL East D-wall Top connection where 3 sampling units were to be selected, the top 3 D-wall panels sorted out of 27 panels according to the values of the randomly generated numbers would be chosen as the panels to be opened up, and the 4th D-wall panel in the sorted list would replace any one of the top 3 originally selected D-wall panels if exposure of coupler connections was found to be not feasible. Likewise, the panel with the highest value out of 10 random numbers generated for EWL West D-wall Top connection was selected to be opened up, and the next panel would replace the originally selected panel if difficulties were encountered.
- 2.3.7 While the lengths of panels range from 2.8m to 7.2m and that the size of the opening up area was about 400mm width for yielding 3 coupler connections in the same layer, it was necessary to determine the exact location of the opening up area on plan at each of the D-wall panels selected as described above. To achieve this, another random number with 5 decimal places valued from 0 to 1 was generated from a uniform distribution for each of the selected D-wall panels and the random number would be multiplied by the corresponding D-wall panel length in order to determine the distance of a reference point from the southern end of the D-wall panel. The reference point will be used as the basis for locating the sampling unit on plan. **Figure 1** illustrates the determination of the reference point.

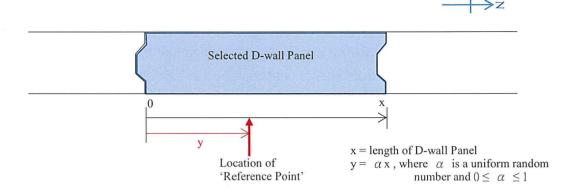


Figure 1 – Determine "Reference Point" for locating the sampling unit on plan

- 2.3.8 The exact location of a sampling unit, which should yield 3 coupler connections at the same layer, at each selected D-wall panel was determined based on the location of the reference point using the "best compliant" approach. The "best compliant" approach provided that once a reference point at the selected D-wall panel was determined, unless the reference point could not physically be accessible or opened up due to physical constraints, all reasonable efforts—should be made to obtain the data at such selected location. If it was impossible to expose such coupler connections, the 3 coupler connections which were at the nearest location to the reference point would be chosen as the sampling unit instead. And if the nearest location with 3 coupler connections measured from the Phase 1 selected location were of equal distance on both sides, then the north side from the Phase 1 selected location would be chosen for opening up.
- 2.3.9 While the determination of reference point locations was carried out in advance, the exact location of each sampling unit was determined on site by Ground Penetration Radar (GPR).
- 2.3.10 To maintain the independence and impartiality of the random selection process, the generation of random numbers, the sorting of D-wall panels in numerical orders, and the calculations of reference point locations were under the full control of the HKU Statistics Team.

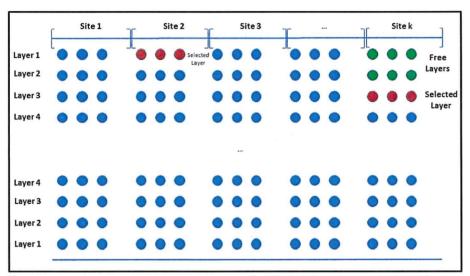
Phase 2 sampling selection

2.3.11 Phase 2 sampling selection was to determine the layer of coupler

connections to be exposed for workmanship verification at the locations selected in Phase 1. According to the drawings and construction records, there could be at most 5 layers of coupler connections at the top or soffit of the slab at the D-wall to slab connections, which varies from panel to panel. In view of the uncertainty in the number of layers of coupler connections existing at the locations selected in Phase 1, a random permutation of the numbers "1" to "5" was generated for each panel selected in Phase 1. The layer of coupler connections to be exposed as the sampling unit was prioritized according to such permutation. The details are described in the following paragraphs.

- 2.3.12 For each panel selected in Phase 1 sampling selection, a numerical sequence consisting of numbers from "1" to "5" was randomly generated. The numerical sequence of each panel represented the order of priority of the layer of coupler connections to be exposed for that panel at the selected location. In parallel, MTRCL reviewed the approved drawings and as-constructed records and advised the maximum number of layers of coupler connection present in the selected locations.
- 2.3.13 At each selected location, the maximum number of layers of coupler connections advised by MTRCL was then mapped with the numerical sequence generated by the HKU Statistics Team. The first number in the numerical sequence that was smaller than or equal to the maximum number of layers of coupler connections at the selected location was chosen as the layer to be exposed. For example, if a numerical sequence of "4, 2, 5, 1, 3" was generated for a particular panel and that such panel was revealed to have at most 3 layers of coupler connections, then the sampling unit for that panel would be located in the 2nd layer of coupler connections counting from the outermost layer because the first number "4" was greater than the maximum number of layers of coupler connections and the subsequent number "2" was smaller than the number of actual layers of coupler connections, making such layer possible to be exposed.
- 2.3.14 The procedures described in paragraphs 2.3.12 and 2.3.13 were repeated for all panels selected in Phase 1 until the layers of coupler connections at which the sampling units at all the selected panels were determined.
- 2.3.15 It is expected that each layer will yield 3 coupler connections. If 3

couplers at Layer 3 are selected, the 6 coupler connections in Layers 1 and 2 will also be examined and included as extra samples (**Figure 2**).



Note: Couplers above the selected layer will be exposed and included as extra samples for verification of workmanship quality.

Figure 2 – An illustration of the proposed sampling scheme with Layer 1 of Site 2 and Layer 3 of Site k being selected

2.4 Samples selection meetings

2.4.1 Two meetings were held between the Government and MTRCL for the random selection of sampling units at EWL slab and NSL slab for Purpose (ii) investigation.

First samples selection meeting

- 2.4.2 The first meeting was held on 5 December 2018 at 11:00 am at Run Run Shaw Building, The University of Hong Kong. The aim of the first meeting was to conduct the Phase 1 sampling selection exercise for EWL slab.
- 2.4.3 The random selection was conducted by the HKU Statistics Team and witnessed by representatives of the Government and MTRCL.
- 2.4.4 The HKU Statistics Team briefed the meeting the random sampling arrangement described in section 2.3 above. The Government and MTRCL also agreed on the sampling population described in section 2.2 above. Further, the meeting discussed and agreed the "best compliant" approach described in section 2.3.8 above.

Second samples selection meeting

- 2.4.5 The second meeting was held on 10 December 2018 at 10:00 am at Run Run Shaw Building, The University of Hong Kong. The aim of the second meeting was to conduct the Phase 1 sampling selection for NSL slab, and the Phase 2 sampling selection for both EWL slab and NSL slab.
- 2.4.6 Again the random selection was conducted independently by the HKU Statistics Team and witnessed by representatives of the Government and MTRCL.
- 2.4.7 At the beginning of this second meeting, MTRCL's representative advised that after checking the locations of the sampling units at the EWL slab selected in the first sample selection meeting against the as-constructed records, the following difficulties were envisaged in exposing the coupler connections at certain selected locations. After discussion, the solutions as stated below were agreed among the parties.

Location	Difficulty encountered	Agreed solution
selected in		
first meeting		
EWL West	Tie-beam was found at the	Apply "best compliant"
D-wall Top –	selected reference point at offset	approach to exposing the
Panel WH35	3.416m, and there was no rebar	nearest coupler
	connecting D-wall and EWL	connections from the
	slab at that offset location.	selected reference point.
EWL East	Mass concrete of at least 1 metre	Replace these three
D-wall Soffit	thick has been cast against the	locations with the first
- Panels	soffit of EWL slab at these three	three panels on the
EH27, EH29	panels of East D-wall, which	"waiting list" produced
and EH38	obstructs the exposure of coupler	at the first meeting under
	connections.	the Group (A3) - EWL
		East D-wall Soffit
		connection (i.e. EH107,
		EM90 and EH97)
EWL West	Mass concrete of at least 1 metre	Replace these three
D-wall Soffit	thick has been cast against the	locations with the first
- Panels	soffit of EWL slab at these three	three panels on the
WH30,	panels of West D-wall, which	"waiting list" produced
WH36 and	obstructs the exposure of coupler	at the first meeting under
WH42	connections.	the Group (A4) - EWL
		West D-wall Soffit
		connection (i.e. WM78,
		WH68 and WM133)

2.4.8 After all the locations to be opened up and the respective layers of coupler connections to be exposed were selected, the random sampling results were validated at the meeting to ensure that all parties agreed with the sampling results.

3. Statistical analysis of the PAUT results obtained from the opening up investigation

3.1 Verification of defective rates

- 3.1.1 The opening up exercise took place from December 2018 to April 2019. Throughout the period, I noted that the opening up and PAUT results were published and regularly updated on the Highways Department's website.
- 3.1.2 After all the PAUT results became available, I was invited to verify the accuracy of the estimated defective rate calculated on the basis of the PAUT results provided by MTRCL.
- 3.1.3 The opening up exercises were carried out in the EWL and NSL slabs independently. The target sample size in each slab was at least 84 as suggested. MTRCL provided 90 valid PAUT results for EWL slab of which 25 were found to be defective, and 93 valid PAUT results for NSL slab of which 23 were found to be defective. I reviewed the opening up results, and found no strong statistical evidence of clustering in the sample.
- 3.1.4 Using the exact binomial formula as listed in paragraph 1.4.1 above with a 95% confident level, the upper bounds were estimated to be 36.6% and 33.2% for EWL and NSL slabs respectively.
- 3.1.5 I noted that MTRCL used the Clopper-Pearson method [OU6/9684 9685] to calculate the upper bounds of the defective rates for EWL and NSL slabs and also arrived at 36.6% and 33.2% respectively. I considered that the methods and calculations performed by MTRCL were appropriate and in order.

3.2 The issue of partial engagement of coupler connections

3.2.1 The current analysis of the quality of workmanship of coupler

connections is based on binomial analysis. I have not received any instructions to review and comment on any proposals regarding partial engagement of coupler connections and the associated statistical analysis. As far as I understand, some tests on the partially engaged coupler connections were carried out by MTRCL. However, these test results failed to comply with all the engineering requirements and there was no proposal received by the Government regarding the use of any residual strength that partially engaged coupler connections may provide.

- 3.2.2 In the design stage of the Holistic Proposal, I verified the suggestion using a binomial analysis by MTRCL. I considered the binomial analysis appropriate because it uses the minimum number of assumptions. From the statistical perspective, the fewer assumptions one makes, the more desirable is the statistical analysis. More assumptions may introduce more uncertainty as some assumptions cannot be verified easily. If the assumptions made are not entirely true, the conclusion drawn from the statistical analysis may no longer be valid.
- 3.2.3 In the first part of the Inquiry, there were discussions regarding the residual strength of the partially engaged coupler connections. coupler connections with insufficient engagement can be allowed and taken into account in the design, multinomial analysis may be relevant. However, to design a multinomial analysis procedure, one needs to classify a failed (i.e. partially engaged) coupler connection into several classes with different ranges of engagement lengths. determine the number of classes, the range covered in each class, and the corresponding allowable residual strength of the coupler connections, if any, in each class. However, I am not aware of any reference standards which may facilitate or allow for such determination. Moreover, a testing plan needs to be developed to provide a reliable and representative estimate for the allowable residual strength, if any, in each For each class, there should be a sufficient number of samples so as to achieve a reasonably small margin of error with a high level of confidence. That means more classes would necessitate a much greater sample size as compared to the binomial analysis.
- 3.2.4 As mentioned above, there are many additional arbitrary decisions which would need to be made for a multinomial analysis such as the number of classes, the range of engagement lengths of each class and so on.

Different combinations of these decisions will end up with different estimates. For example, consider the case with 3 classes, namely (A) pass, (B) partial pass, and (C) fail. Consider two designs with different ranges for classes (B) and (C) with the first design having 70% (A), 25% (B) and 5% (C) and the second design having 70% (A), 15% (B) and 15% (C). The estimates to be arrived at based on the two designs can be very different which would introduce room for manipulation of the data.

- 3.2.5 I consider the use of binomial analysis reasonable as it involves less arbitrary decisions in the design. It is also more practical due to the smaller sample size required and hence less cost and time implications. It is a more suitable approach for achieving the primary objective to assess whether the quality of coupler connections is in compliance with specification or not.
- 4. Statistical analysis to derive a combined defective rate to account for the condition on both sides of coupler connections at locations where EWL slab was connected to the D-wall via capping beam

4.1 The issue

- 4.1.1 I was given to understand that some of the randomly selected panels to be opened up are with a different configuration from the others in that there is a capping beam resting on top of the D-wall such that the coupler connection is located within the EWL slab instead of partly embedded in the D-wall. During the opening up exercise, some of the coupler connections on the side of the capping beam, in addition to the side of the platform slab, were exposed as well. As a result, the workmanship of coupler connections on the capping beam side also came to light.
- 4.1.2 Based on the information provided by MTRCL, out of the 11 exposed coupler connections on the capping beam side, 2 of them were found to be not in compliance with the manufacturer's specification, i.e. with more than two threads exposed. On the platform slab side of the same panels, 2 out of 7 coupler connections with valid PAUT results are found to be not in compliance with the manufacturer's specification. This finding has brought up a new situation on the acceptance criteria of the coupler connection. While a coupler connection could only perform as

intended when the rebars on both sides of the coupler are properly screwed in, it is necessary to consider the workmanship of the coupler connection of both sides. Only in the situation where the connections on both sides are proper can a coupler connection be considered as satisfactory for this type of configuration. Failure in either side or both sides of the coupler connection will result in a defective coupler connection as a whole. It is therefore necessary to find a way to take into account the failure rates on both sides of the coupler connections for those EWL panels with capping beam. This situation arose due to some unexpected observations, and was not contemplated when designing the sampling plan. I was therefore requested to review the situation and make a suggestion on how the effect of defective coupler connections on both sides of a coupler could be reflected in a single defective rate, namely in statistical terms.

4.2 Derivation of a combined defective rate

- 4.2.1 As mentioned above, a satisfactory coupler connection is defined as one for which both sides of the connection satisfy the installation requirements. Failure in either side or both sides of the coupler connection will result in a defective coupler connection.
- 4.2.2 To assess the combined effect of the defective workmanship on both sides of a coupler connection, a two-step approach is used. Based on the information provided by MTRCL, the proportions of failed coupler connections on each sides could be calculated. Assuming that the quality of workmanship on the two sides of the coupler connection were independent, a combined defective rate taking into account the failed proportions of both sides of the coupler connection can be computed using simple probability theory described as follows.
- 4.2.3 Let p_B be the proportion of defective coupler connections for those panels with capping beam at EWL level. Assume that the quality of workmanship on the two sides are independent. Since the coupler connection is considered satisfactory if and only if the connection on both sides are satisfactory, therefore it is easy to see that

$$1 - p_B = (1 - p_{B1})(1 - p_{B2})$$

where p_{B1} and p_{B2} are the proportions of unsatisfactory connections on the platform slab side and capping beam side respectively. The proportion (i.e. the combined defective rate) can be estimated by

$$\hat{p}_B = 1 - (1 - \hat{p}_{B1})(1 - \hat{p}_{B2}).$$

According to the data provided by MTRCL, we have $\hat{p}_{B1} = \frac{2}{7}$ and $\hat{p}_{B2} = \frac{2}{11}$, therefore p_B is estimated to be

$$\hat{p}_B = 1 - \left(1 - \frac{2}{7}\right) \left(1 - \frac{2}{11}\right) \approx 0.4156 = 41.56\%$$
.

- 4.2.4 The above combined defective rate only represents the defective rate at the opened up locations. To determine the defective rate applicable for all panels of similar configurations at 95% confidence level, a statistical inference is required.
- 4.2.5 Using the delta method and after some algebraic manipulation, the variance of \hat{p}_B is given by

$$\operatorname{Var}(\hat{p}_B) = \{ (1 - \hat{p}_{B1})(1 - \hat{p}_{B2}) \}^2 \left[\frac{\hat{p}_{B1}}{7(1 - \hat{p}_{B1})} + \frac{\hat{p}_{B2}}{11(1 - \hat{p}_{B2})} \right] = 0.0264.$$

Using the normal approximation, the upper bound of a one-sided 95% confidence interval for p_B is given by

$$0.4156 + 1.645 \times \sqrt{0.0264} = 0.6829.$$

4.2.6 The above calculation represents construction of a one-sided 95% confidence interval with reference to the proportion of the number of opening up locations against the actual number of panels with the same configuration. And the upper bound value of the 95% confidence interval for the combined defective rate using the normal approximation was found to be 68.3%.

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POSITIONS 2018 – present	Patrick S. C. Poon Professor in Statistics and Actuarial Science, University of Hong Kong
2017 – present	Head of Department Department of Statistics and Actuarial Science University of Hong Kong, Hong Kong
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RESEARCH INTERESTS	Clinical trial methodology and adaptive design Data mining, machine learning, AI Bayesian methods and inference Survival analysis Big data and high-dimensional data analysis

C/C++, Python, SAS, R, Matlab, WinBUGS, East, PASS, nQuery

COMPUTER

SKILLS

PROFESSIONAL

MEMBERSHIP American Statistical Association (ASA)

Eastern North American Region of the Biometric Society (ENAR)

International Society for Bayesian Analysis (ISBA)

Institute of Mathematical Statistics (IMS)

International Chinese Statistical Association (ICSA)

HONORS & AWARDS

2013 Fellow of the American Statistical Association

2012 Elected Member of the International Statistical Institute

2009 James E. Grizzle Distinguished Alumni Award, Department of Biostatis-

tics, University of North Carolina at Chapel Hill.

STUDENTS ADVISED

Over 30 M.S., M.Phil., Ph.D., Post-doctor fellows, and Research Assistants/Associates

PRESENTATIONS

Over 200 invited talks, presentations, or seminars in conferences and universities.

GRANTS

Multiple grants from Hong Kong Research Grants Council and National Cancer Institute.

EDITORIAL SERVICES

Associate Editor Statistical Analysis and Data Mining (2018 – present)

Associate Editor Japanese Journal of Statistics and Data Science (2018 – present)

Associate Editor Journal of the American Statistical Association (2013 – present)

Associate Editor Contemporary Clinical Trials (2012 – present)

Associate Editor Bayesian Analysis (2009 – 2015)

Guest Editor Special issue on Clinical Trials for Personalized Medicine in Contempo-

rary Clinical Trials (2012)

Grant Reviewer Hong Kong Research Grants Council

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Book Reviewer John Wiley & Sons, CRC Press (Taylor & Francis Group), Springer

PUBLICATIONS

Book

- 1. Yin, G. (2012). Clinical Trial Design: Bayesian and Frequentist Adaptive Methods by John Wiley & Sons (Wiley Series in Probability and Statistics). Hoboken, New Jersey, USA. Japanese translated version by Teramukai, S. and Daimon, T. (2014), Medical Publications, Tokyo, Japan.
- 2. Yin, G. and Shi, H. (2018). Statistical Design and Analysis in Clinical Trials (in Chinese). Higher Education Press, China.

Papers under Review/Revision

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- 3. Yang, Z. and **Yin**, **G**.* (2018). Fractional design for late-onset toxicities in oncology dose-finding studies. *CCR*.
- 4. Yang, Z. and Yin, G.* (2019). An alternative approach for estimating the number needed to treat for survival endpoints. *Annals of Internal Medicine*.
- 5. Zhang, C. and **Yin**, **G**.* Re-analysis of the data of laser peripheral iridotomy for the prevention of angle closure. *The Lancet*.
- 6. Jiang, F. and Yin, G. (2018). Bayesian reflected non-local priors with application to edge detection in drone images. Submitted to IEEE Transactions on Image Processing.
- 7. Jiang, F., Yin, G. and Dominici, F. (2019). Bayesian model selection approach to multiple change-points detection with non-local prior distributions. Submitted to IEEE Transactions on Pattern Analysis and Machine Intelligence.
- 8. Yang, Z., Lin, Y., **Yin**, **G.** and Yuan, Y. (2018). Sample size re-estimation in adaptive enrichment strategies. *Submitted to Clinical Trials*.
- 9. Ma, D., Liu, B., Cao, D., and **Yin, G.** (2018). A coarse-to-fine framework for music generation. Submitted to IJCAI.
- 10. Liu, B. and Yin, G. (2018). Air quality forecasting with convolutional LSTM. Submitted to IJCAI.
- 11. He, B., Liu, Y., Wu, Y. and **Yin, G.** (2018). Doubly divided the massive data for prediction using model aggregation. *Submitted to JASA*.
- 12. He, B., Liu, Y., Wu, Y., Yin, G., and Zhao, X. (2018). Functional martingale residual process for high-dimensional Cox regression with model averaging. Submitted to Biometrika.
- 13. Liu, Y. and Yin, G. (2018). Ensemble Delaunay triangulation learner. Submitted to CSDA.
- 14. Liu, Y. and Yin, G. (2018). Nonparametric functional approximation with Delaunay triangulation. Submitted to JCGS.

- 15. Liu, Y. and Yin, G. (2018). Collaborative gradient boosting. Submitted to Nuerocomputing.
- 16. Lin, R., Yuan, Y., **Yin, G.** and Thall, P. (2018). Bayesian hierarchical random-effects metaanalysis and design of phase I clinical trials. *Submitted to Biometrics*.
- 17. Lin, R. and Yin, G.* (2018). Adaptive model selection design for identifying optimal biological dose in phase I/II clinical trials. Submitted to Bayesian Analysis.
- 18. Lam, C. K. and **Yin**, **G.*** (2018). A variable selection approach to multiple change-points detection. Submitted to Statistics and Its Interface.

Proceedings of Top Conferences in AI and Machine Learning

- 19. Gu, J. and Yin, G.* (2019). Fast algorithm for generalized multinomial models with ranking data. *ICML* 2019 (36th International Conference on Machine Learning).
- 20. Zhang, C. and Yin, G.* (2019). Fast and stable maximum likelihood estimation for incomplete multinomial model. *ICML* 2019 (36th International Conference on Machine Learning).
- 21. Jiang, F., Yin, G. and Dominici, F. (2018). Bayesian model selection approach to boundary detection with non-local priors. Proceeding of the 32nd Conference on Neural Information Processing Systems (NeurIPS 2018), Montrèal, Canada.

Statistics in Top Medical Journals (corresponding author*)

- 22. Yin, G.* and Zhang C. (2019). Reanalysis of the data comparing prophylactic cranial irradiation vs observation in patients with locally advanced non-small cell lung cancer. *Journal of American Medical Association Oncology*. In press.
- 23. Yin, G.* and McCaw, Z. R. (2019). Design of non-inferiority trials for hypofractionated vs conventional radiotherapy among cancer patients. *Journal of American Medical Association Oncology*. In press.
- 24. McCaw, Z. R., Yin, G. and Wei, L. J. (2019). Interpreting the treatment effect of new-generation drug-eluting stents. *Circulation*. In press.
- 25. Yang, Z. and Yin, G.* (2019). The number needed to treat for both binary and survival endpoints. *British Medical Journal*.
- 26. Yang, Z., McCaw, Z. R. and **Yin, G.*** (2019). Reanalysis of the effectiveness of caplacizum-ab treatment for acquired thrombotic thrombocytopenic purpura. *New England Journal of Medicine*. In press.
- 27. Yang, Z., McCaw, Z. R. and Yin, G.* (2019). Interpreting the long-term benefit of radical prostatectomy in prostate cancer. *New England Journal of Medicine* 380, 1083–1084.
- 28. Morita, S., Sakamaki, K., and **Yin, G.** (2015). Detecting overall survival benefit derived from survival postprogression rather than progression-free survival. *Journal of the National Cancer Institute* **107** (8), djv133, doi:10.1093/jnci/djv133.

- 29. Lee, J. J., Chen, N., and Yin, G. (2012). Worth adapting? Revisiting the usefulness of outcome-adaptive randomization. *Clinical Cancer Research* 18, 4498–4507.
- 30. Yuan, Y. and Yin, G. (2011). On the usefulness of outcome-adaptive randomization. *Journal* of Clinical Oncology 29, e390–e392.

Statistical Methodologies

- 31. Yin, G.* and Shi, H. (2019). Demystify Lindley's paradox: A new interpretation of p-value as the posterior probability of the null. Submitted to International Statistical Review.
- 32. Shi, H. and Yin, G.* (2019). P-value: A blessing or a curse for evidence-based studies? Submitted to The American Statistician.
- 33. Shi, H., Zhang, T. and **Yin**, **G.*** (2019). START: Single-to-double arm transition design for phase II clinical trials. *Submitted to Pharmaceutical Statistics*.
- 34. Wang, N. and Yin, G. (2019). Convergence rates of the blocked Gibbs sampler with random scan in the Wasserstein metric. *Stochastics*. In press.
- 35. Yan, X., Yin, G. and Zhao, X. (2019). Subgroup analysis in censored linear regression. *Statistica Sinica*. In press.
- 36. Jiang, F., Cheng, Q., Yin, G. and Shen, H. (2019). Functional censored quantile regression. Journal of the American Statistical Association. In press.
- 37. Wang, Y., Zou, C., Yin, G.* and Wang, Z. (2018). Multiple change-points detection in high dimension. *Revised for Statistica Sinica*.
- 38. Lam, C. K., Lin, R. and Yin, G.* (2018). Nonparametric overdose control for dose finding in drug-combination trials. *Journal of Royal Statistical Society C Applied Statistics*. In press.
- 39. Shi, H. and **Yin**, **G.*** (2019). Control of type I error rates in Bayesian sequential designs. *Bayesian Analysis* 14, 399–425.
- 40. Zheng, S, Lin, R., Guo, J. and **Yin, G.** (2019). Testing homogeneity of high-dimensional covariance matrices. *Statistica Sinica*, in press.
- 41. Lin, R. and Yin, G.* (2018). Uniformly most powerful Bayesian interval design for dose finding. *Pharmaceutical Statistics* 17, 710–724.
- 42. Lam, C. K., Xu, Y., and Yin, G. (2018). Dynamic portfolio choice without cash. *Quantitative Finance* 19, 313–326.
- 43. Liu, Y. and **Yin**, **G**. (2018). Average holding price. *Annals of Financial Economics* **13**, No. 01. https://doi.org/10.1142/S2010495218500021
- 44. Shi, H. and Yin, G.* (2018). Bayesian enhancement two-stage design for single-arm phase II clinical trials with binary and time-to-event endpoints. *Biometrics* 74, 1055–1064.
- 45. Jiang, F., Ma, Y., and **Yin, G.** (2018). Kernel-based adaptive randomization toward balance in continuous and discrete covariates. *Statistica Sinica*, in press.

- 46. Li, H., Cao, Z., and **Yin**, **G.*** (2018). Varying-association copula models for multivariate survival data. *Canadian Journal of Statistics*, in press.
- 47. Dong, F., and Yin, G. (2018). Maximum likelihood estimation for incomplete multinomial data via the weaver algorithm. *Statistics and Computing* 28, 1095–1117.
- 48. Yin, G., Chen, N., and Lee, J. J. (2018). Bayesian adaptive randomization and trial monitoring with predictive probability for time-to-event endpoint. *Statistics in Bioscience* 10, pp 420–438.
- 49. Wang, G., Zou, C., and Yin, G. (2018). Change-point detection in multinomial data with a large number of categories. *Annals of Statistics* 46, 2020–2044.
- 50. Shi, H. and Yin, G.* (2018). Boosting conditional logit model. *Journal of Choice Modeling* 26, 48–63.
- 51. Zhang, J., Yin, G., Liu, Y., and Wu, Y. (2018). Censored cumulative residual independent screening for ultrahigh-dimensional survival data. *Lifetime Data Analysis* 24, 273–292.
- 52. Shi, H. and Yin, G.* (2018). Two-stage seamless transition design from open-label single-arm to randomized double-arm clinical trials. *Statistical Methods in Medical Research* 27, 158–171.
- 53. Lin, R. and **Yin**, **G.*** (2017). STEIN: A simple toxicity and efficacy interval design for seamless phase I/II clinical trials. *Statistics in Medicine* **36**, 4106–4120.
- 54. Duan, X., and Yin, G.* (2017). Ensemble approaches to estimating the population mean with missing response. Scandinavian Journal of Statistics 44, 899–917.
- 55. Yin, G.*, Lam, C. K., and Shi, H. (2017). Bayesian randomized clinical trials: from fixed to adaptive design. *Contemporary Clinical Trials* 59, 77–86.
- 56. Shi, H. and Yin, G.* (2017). Landmark cure rate models with with time-dependent covariates. Statistical Methods in Medical Research 26, 2042–2054.
- 57. Liu, Y. and Yin, G.* (2017). Partitioned log-rank tests for the overall homogeneity of hazard rate functions. *Lifetime Data Analysis* 23, 400–425.
- 58. Lin, R. and Yin, G.* (2017). Bayesian optimal interval design for dose finding in drug-combination trials. Statistical Methods in Medical Research 26, 2155–2167.
- 59. Zhao, X., Wu, Y., and **Yin, G.** (2017). Sieve maximum likelihood estimation for a general class of accelerated hazards models with bundled parameters. *Bernoulli* **23**, 3385–3411.
- 60. Lin, R. and **Yin**, **G**.* (2017). Nonparametric overdose control with late-onset toxicity in phase I clinical trials. *Biostatistics* **18**, 180–194.
- 61. Wu, Y. and Yin, G.* (2017). Cure rate quantile regression accommodating both finite and infinite survival times. *Canadian Journal of Statistics* 45, 29–43.
- 62. Wu, Y. and Yin, G.* (2017). Multiple imputation for cure rate quantile regression with censored data. *Biometrics* 73, 94–103.
- 63. Shi, H. and **Yin**, **G.*** (2017). Bayesian two-stage design for phase II clinical trials with switching hypothesis tests. *Bayesian Analysis* **12**, 31–51.

- 64. Lin, R. and **Yin**, **G**.* (2016). Bayesian optimal interval design for drug-combination trials. Frontiers of Biostatistical Methods and Applications in Clinical Oncology. Springer.
- 65. Lin, R. and Yin, G.* (2016). Robust optimal interval design for high-dimensional dose finding in multi-agent combination trials. ICSA book proceedings for the symposium at Calgary. Springer.
- 66. Lin, R., Liu, Z., Zheng, S., and Yin, G. (2016). Power computation for hypothesis testing with high-dimensional covariance matrices. *Computational Statistics & Data Analysis* 104, 10–23.
- 67. Lin, R. and Yin, G.* (2016). Bootstrap aggregating continual reassessment method for dose finding in drug-combination trials. *Annals of Applied Statistics* **10**, 2349–2376.
- 68. Li, H., Duan, X., and **Yin**, **G.*** (2016). Generalized method of moments for additive hazards model with clustered dental survival data. *Scandinavian Journal of Statistics* **43**, 1124–1139.
- 69. Xu, J., Yin, G.*, Ohlssen, D., and Bretz, F. (2016). Bayesian two-stage dose finding for cytostatic agents via model adaptation. *Journal of Royal Statistical Society C Applied Statistics* 65, 465–482.
- 70. Lin, R. and Yin, G.* (2015). Bayes factor and posterior probability: complementary statistical evidence to p-value. *Contemporary Clinical Trials* 44, 33–35.
- 71. Ro, K., Zou, C., Wang, Z., and Yin, G. (2015). Outlier detection for high dimensional data. *Biometrika* 102, 589–599.
- 72. Jin, I. H., Huo, L., Yuan, Y., and **Yin, G.** (2015). Phase I trial design for drug combinations with Bayesian model averaging. *Pharmaceutical Statistics* **14**, 108–119.
- 73. Yin, G. and Lin, R. (2015). Continual reassessment methods. *Modern Approaches to Clinical Trials Using SAS: Classical, Adaptive, and Bayesian Methods* Chapter 5, edited by R. Zink and S. Menon, pp. 133–162. SAS Institute Inc., Cary, NC, USA.
- 74. Lin, R. and Yin, G.* (2015). Overview of adaptive randomization. *Modern Approaches to Clinical Trials Using SAS: Classical, Adaptive, and Bayesian Methods* Chapter 9, edited by R. Zink and S. Menon, pp. 247–272. SAS Institute Inc., Cary, NC, USA.
- 75. Lin, R. and Yin, G.* (2015). Sample size re-estimation in adaptively randomized clinical trials with missing data. *Modern Adaptive Randomized Clinical Trials: Statistical and Practical Aspects*, edited by O. Sverdlov, pp. 269–286. Chapman & Hall/CRC Press.
- 76. Wu, Y., Ma, Y., and **Yin, G.*** (2015). Smoothed and corrected score approach to censored quantile regression with measurement errors. *Journal of the American Statistical Association* **110**, 1670–1683.
- 77. **Yin**, **G.** and Lin, R. (2015). Comments on 'Competing designs for drug combination in phase I dose-finding clinical trials' by M-K. Riviere, F. Dubois and S. Zohar *Letter to Editor Statistics in Medicine* **34**, 13–17.
- 78. Wu, Y. and Yin, G.* (2015). Conditional quantile screening in ultrahigh-dimensional heterogeneous data. *Biometrika* 102, 65–76.

- 79. Wang, X., Zhang, J., Yu, L., and **Yin**, **G.*** (2014). Generalized partially linear single-index model for zero-inflated count data. *Statistics in Medicine* **34**, 876–886.
- 80. Zou, C., Yin, G.*, Feng, L., and Wang, Z. (2014). Nonparametric maximum likelihood approach to multiple change-point problems. *Annals of Statistics* 42, 970–1002.
- 81. Xu, J. and Yin, G.* (2014). Two-stage adaptive randomization for delayed response in clinical trials. Journal of Royal Statistical Society C Applied Statistics 63, 559–578.
- 82. Yin, G., Zeng, D., and Li, H. (2014). Censored quantile regression with varying coefficients. Statistica Sinica 24, 855–870.
- 83. Wu, Y. and Yin, G.* (2013). Cure rate quantile regression for censored data with a survival fraction. *Journal of the American Statistical Association* 108, 1517–1531.
- 84. Liu, S., Yin, G., and Yuan, Y. (2013). Bayesian data augmentation dose finding with continual reassessment method and delayed toxicity. *Annals of Applied Statistics* 7, 2138–2156.
- 85. Yin, G. and Ma, Y. (2013). Pearson-type goodness-of-fit test with bootstrap maximum likelihood estimation. *Electronic Journal of Statistics* 7, 412–427.
- 86. Shi, Y. and **Yin**, **G.*** (2013). Escalation with overdose control for phase I drug-combination trials. *Statistics in Medicine* **32**, 4400–4412.
- 87. Ma, Y. and Yin, G.* (2013). Testing overall and subpopulation treatment effects with measurement errors. *Statistica Sinica* 23, 1019–1042.
- 88. Yin, G., Zheng, S., and Xu, J. (2013). Two-stage dose finding for cytostatic agents in phase I clinical trials. *Statistics in Medicine* **32**, 644–660.
- 89. Yin, G., Zheng, S., and Xu, J. (2013). Fractional dose-finding methods with late-onset toxicity in phase I clinical trials. *Journal of Biopharmaceutical Statistics* 23, 856–870.
- 90. Gu, X., Yin, G., and Lee, J. J. (2013). Bayesian two-stage Lasso strategies for biomarker selection in personalized medicine development. *Contemporary Clinical Trials* 36, 642–650.
- 91. Huo, L., Yuan, Y., and **Yin, G.*** (2012). Bayesian dose finding for combined drugs with discrete and continuous doses. *Bayesian Analysis* 7, 235–252.
- 92. Diao, G. and Yin, G. (2012). A general transformation class of semiparametric cure rate frailty models. *Annals of the Institute of Statistical Mathematics* 64, 959–989.
- 93. Yin, G., Chen, N., and Lee, J. J. (2012). Phase II trial design with Bayesian adaptive randomization and predictive probability. *Journal of Royal Statistical Society C Applied Statistics* 61, 219–235.
- 94. Garcia, T., Ma, Y., and **Yin**, **G.** (2011). Efficiency improvement in a class of survival models through model-free covariate incorporation. *Lifetime Data Analysis* **7**, 552–565.
- 95. Yuan, Y. and **Yin**, **G.** (2011). Dose-response curve estimation: A semiparametric mixture approach. *Biometrics* **67**, 1543–1554.

- 96. Yuan, Y. and **Yin**, **G.** (2011). Robust EM continual reassessment method in oncology dose finding. *Journal of the American Statistical Association* **106**, 818–831.
- 97. Yin, G., Ma, Y., Liang, F., and Yuan, Y. (2011). Stochastic generalized method of moments. Journal of Computational and Graphical Statistics 20, 714–727.
- 98. Yuan, Y. and **Yin**, **G.** (2011). Bayesian hybrid dose-finding design in phase I oncology clinical trials. *Statistics in Medicine* **30**, 2098–2108.
- 99. Yuan, Y. and **Yin**, **G.*** (2011). Bayesian phase I/II adaptively randomized oncology trials with combined drugs. *Annals of Applied Statistics* 5, 924–942.
- 100. Ma, Y. and **Yin**, **G**. (2011). Censored quantile regression with covariate measurement errors. *Statistica Sinica* **21**, 949–971.
- 101. Lei, X., Yuan, Y., and **Yin, G.*** (2011). Bayesian phase II clinical trial design with time-to-event adaptive randomization. *Lifetime Data Analysis* 17, 156–174.
- 102. **Yin, G.** and Yuan, Y. (2010). Correspondence to the discussion of "Bayesian dose finding in oncology for drug combinations by copula regression." *Journal of Royal Statistical Society C Applied Statistics* **59**, 544–546.
- 103. Ma, Y. and Yin, G.* (2010). Semiparametric median residual life model and inference. Canadian Journal of Statistics 38, 665–679.
- 104. Yuan, Y. and Yin, G. (2010). Bayesian quantile regression for longitudinal studies with non-ignorable missing data. *Biometrics* 66, 105–114.
- 105. Yin, G. and Yuan, Y. (2010). Bayesian approach for adaptive design. Handbook of Adaptive Designs in Pharmaceutical and Clinical Development, A. Pong, and S.-C. Chow, (eds.), Chapter 3. pp. (3-1)-(3-19). Boca Raton, FL: Chapman & Hall/CRC.
- 106. Yin, G. and Nieto-Barajas, L. E. (2009). Bayesian cure rate model accommodating multiplicative and additive covariates. *Statistics and Its Interface* 2, 513–521.
- 107. Yin, G. and Li, H. (2009). Least squares estimation of varying-coefficient hazard regression with application to breast cancer dose-intensity data. *Canadian Journal of Statistics* 37, 659–674.
- 108. Yin, G. and Yuan, Y. (2009). Bayesian model averaging continual reassessment method in phase I clinical trials. *Journal of the American Statistical Association* 104, 954–968.
- 109. Yin, G. (2009). Bayesian generalized method of moments (with discussion). Bayesian Analysis 4, 191–208; and Rejoinder, 217–222.
- 110. Yuan, Y. and Yin, G. (2009). Bayesian dose finding by jointly modeling toxicity and efficacy as time-to-event outcomes. *Journal of Royal Statistical Society C Applied Statistics* 58, 719–736.
- 111. Li, H. and Yin, G.* (2009). Generalized method of moments for linear regression with clustered failure time data. *Biometrika* 96, 293–306.
- 112. **Yin, G.** and Yuan, Y. (2009). Bayesian dose finding in oncology for drug combinations by copula regression. *Journal of Royal Statistical Society C Applied Statistics* **58**, 211–224.

- 113. Yin, G. (2009). Bayesian chi-squared goodness-of-fit test for censored data. *Journal of Statistical Planning and Inference* **139**, 1474–1483.
- 114. **Yin, G.** and Yuan, Y. (2009). A latent contingency table approach to dose finding for combinations of two agents. *Biometrics* **65**, 866–875.
- 115. Yin, G. (2008). Bayesian transformation cure frailty models with multivariate failure time data. Statistics in Medicine 27, 5929–5940.
- 116. Yuan, Y. and **Yin**, **G.** (2008). Sequential continual reassessment method for two-dimensional dose finding. *Statistics in Medicine* **27**, 5664–5678.
- 117. Yin, G., Li, H., and Zeng, D. (2008). Partially linear additive hazards regression with varying coefficients. *Journal of the American Statistical Association* 103, 1200–1213.
- 118. Yin, G., Zeng, D., and Li, H. (2008). Power-transformed linear quantile regression with censored data. *Journal of the American Statistical Association* 103, 1214–1224.
- 119. Ma, Y. and Yin, G. (2008). Cure rate model with mismeasured covariates under transformation. *Journal of the American Statistical Association* 103, 743–756.
- 120. Nieto-Barajas, L. E. and Yin, G. (2008). Bayesian semiparametric cure rate model with an unknown threshold. *Scandinavian Journal of Statistics* 35, 540–556.
- 121. Li, H., Yin, G., and Zhou, Y. (2007). Local likelihood with time-varying coefficient additive hazards model. *Canadian Journal of Statistics* 35, 321–337.
- 122. Ji, Y., Yin, G., Tsui, K.-W., Kolonin, M. G., Sun, J., Arap, W., Pasqualini, R., and Do, K.-A. (2007). Bayesian mixture models for complex high-dimensional count data in phage display experiments. *Journal of Royal Statistical Society C Applied Statistics* 56, 1–14.
- 123. Cong, X., Yin, G., and Shen, Y. (2007). Marginal analysis of correlated failure time data with informative cluster sizes. *Biometrics* **63**, 663–672.
- 124. Yin, G. (2007). Model checking for additive hazards model with multivariate survival data. Journal of Multivariate Analysis 98, 1018–1032.
- 125. Ji, Y., Li, Y., and **Yin, G.** (2007). Bayesian dose finding in phase I clinical trials based on a new statistical framework. *Statistica Sinica* 17, 531–547.
- 126. Yin, G. and Ibrahim, J. (2006). Bayesian transformation hazard models. The Second Lehmann Symposium-Optimality, IMS Lecture Notes-Monographs Series, 170–182.
- 127. **Yin, G.**, Li, Y., and Ji, Y. (2006). Bayesian dose-finding in phase I/II trials using toxicity and efficacy odds ratio. *Biometrics* **62**, 777–784.
- 128. Zeng, D., Yin, G., and Ibrahim, J. (2006). Semiparametric transformation models for survival data with a cure fraction. *Journal of the American Statistical Association* 101, 670–684.
- 129. Yin, G. and Zeng, D. (2006). Efficient algorithm for computing maximum likelihood estimates in linear transformation models. *Journal of Computational and Graphical Statistics* 15, 228–245.

- 130. Yin, G. and Ibrahim, J. (2005). Cure rate models: a unified approach. Canadian Journal of Statistics 33, 559–570.
- 131. Zeng, D., Yin, G., and Ibrahim, J. (2005). Inference for a class of transformed hazard models. Journal of the American Statistical Association 100, 1000–1008.
- 132. Yin, G. and Ibrahim, J. (2005). Bayesian frailty models based on Box-Cox transformed hazards. Statistica Sinica 15, 781–794.
- 133. Yin, G. and Shen, Y. (2005). Self-designing trial combining with classical group sequential monitoring. *Journal of Biopharmaceutical Statistics* 15, 667–675.
- 134. Yin, G. (2005). Bayesian cure rate frailty models with application to a root canal therapy study. *Biometrics* 61, 552–558.
- 135. **Yin, G.** and Ibrahim, J. (2005). A general class of Bayesian survival models with zero and non-zero cure fractions. *Biometrics* **61**, 403–412.
- 136. Yin, G. and Shen, Y. (2005). Adaptive design and estimation in randomized clinical trials with correlated observations. *Biometrics* **61**, 362–369.
- 137. Zeng, D., Lin, D. Y., and **Yin, G.** (2005). Maximum likelihood estimation in proportional odds model with random effects. *Journal of the American Statistical Association* **100**, 470–483.
- 138. Yin, G. and Ibrahim, J. (2005). A class of Bayesian shared gamma frailty models with multivariate failure time data. *Biometrics* 61, 209–217.
- 139. Yin, G. and Cai, J. (2005). Quantile regression models with multivariate failure time data. *Biometrics* 61, 152–162.
- 140. Yin, G. and Zeng, D. (2005). Pair chart test for an early survival difference. Lifetime Data Analysis 11, 117–129.
- 141. Yin, G. and Hu, J. (2004). Two simulation methods for constructing confidence bands under the additive risk model. *Journal of Biopharmaceutical Statistics* 14, 389–402.
- 142. Hu, J., Yin, G., Morris, J. S., Zhang, L., and Wright, A. F. (2004). Entropy and survival-based weights to combine Affymetrix array types and analyze differential expression and survival. *Methods of Microarray Data Analysis IV, Critical Assessment of Microarray Data Analysis*, eds. J. S. Shoemaker and S. M. Lin, pp. 95–108.
- 143. Morris, J. S., **Yin**, **G.**, Baggerly, K., Wu, C., and Zhang, L. (2004). Pooling information across different studies and oligonucleotide chip types to identify prognostic genes for lung cancer. *Methods of Microarray Data Analysis IV, Critical Assessment of Microarray Data Analysis*, eds. J. S. Shoemaker and S. M. Lin, pp. 51–66.
- 144. Yin, G. and Cai, J. (2004). Additive hazards model for multivariate failure time data. *Biometrika* 91, 801–818.
- 145. Hu, J. and **Yin**, **G.** (2003). A semiparametric regression model for oligonucleotide arrays. Journal of Modern Applied Statistical Methods **2**, 256–267.

146. Yin, G., Cai, J., and Kim, J. (2003). Quantile inference with multivariate failure time data. *Biometrical Journal* 45, 602–617.

Collaborative Research

- 147. Zhong, Y. J., Wen, Y. F., Wong, H. M., **Yin, G.**, Lin, R. and Yang, S. Y. (2019). Trends and patterns of disparities in burden of lung cancer in the United States, 1974-2015. *Frontiers in Oncology, Thoracic Oncology*. In press.
- 148. Cardó-Vila, M., Marchió, S., Sato, M., Staquicini, F., Bronk, J., **Yin, G.**, Zurita, A., Lee, J. J., Hong, W., Wistuba, I., Arap, W., and Pasqualini, R. (2016). Interleukin-11 receptor is a candidate target for ligand-directed therapy in lung cancer: analysis of clinical samples and BMTP-11 pre-clinical activity. *American Journal of Pathology* **186**, 2162–2170.
- 149. Wen, Y. F., Wong, H. M., Lin, R., Yin, G., and C. P. McGrath (2015). Inter-ethnic/racial facial variations: A systematic review and Bayesian meta-analysis of photogrammetric studies. *Plos One*, DOI:10.1371/journal.pone.0134525
- 150. Arun, B. K., Dhinghra, K., Valero, V., Kau, S.-W., Broglio, K., Booser, D., Guerra, L., Yin, G., Walters, R., Sahin, A., Ibrahim, N., Buzdar, A. U., Frye, D., Sneige, N., Strom, E., Ross, M., Theriault, R., Vadhan-Raj, S., Hortobagyi, G. N. (2011). Randomized trial of dose intensive neoadjuvant chemotherapy with or without G-CSF in locally advanced breast cancer: long-term results. *Oncologist* 16, 1527–1534.
- 151. Richards, K. L., Zhang, B., Sun, M., Dong, W., Churchill, J., Bachinski, L. L., Wilson, C. D., Baggerly, K. A., Yin, G., Hayes, D. N., Wistuba, I. I. and Krahe, R. (2011). Methylation of the candidate biomarker TCF21 is very frequent across a spectrum of early stage non-small cell lung cancers. *Cancer* 117, 606–617.
- 152. Yuan, P., Kadara, H., Behrens, C., Tang, X., Woods, D., Solis, L. M., Huang, J., Spinola, M., Dong, W., Yin, G., Fujimoto, J., Kim, E., Xie, Y., Girard, L., Moran, C., Hong, W. K., Minna, J. D. and Wistuba, I. I. (2010). Sex determining region Y-box 2 (SOX2) is a potential cell-lineage gene highly expressed in the pathogenesis of squamous cell carcinomas of the lung. *Plos One* 5, e9112. doi:10.1371/journal.pone.0009112
- 153. Arun, B. K., Granville, L. A., Yin, G., Middleton, L. P., Dawood, S., Shu, W.-K., Kamal, A., Hsu, L., Hortobagyi, G. N. and Sahin, A. A. (2010). Gluthation-S-transferase-pi (GST-pi) expression in early breast cancer: associated with outcome and response to chemotherapy. *Cancer Investigation* 28, 554–559.
- 154. Sun, M., Behrens, C., Feng, L., Ozburn, N., Tang, X., **Yin, G.**, Komaki, R., Varella-Garcia, M., Hong, W. K., Aldape, K. D. and Wistuba, I. I. (2009). HER family receptor abnormalities in lung cancer brain metastases and corresponding primary tumors. *Clinical Cancer Research* **15**, 4829–4837.
- 155. Rivera, E., Mejia, J., Arun, B., Adinin, R., Walters, R., Abenaa B., A., Broglio, K., **Yin, G.**, Hortobagyi, G. and Valero, V. (2008). Phase III study comparing the use of docetaxel on an every-three-week versus weekly schedule in the treatment of metastatic breast cancer. *Cancer* **112**, 1455–1461.

- 156. Arun, B., Valero, V., Logan, C., Broglio, K., Rivera, E., Brewster, A., **Yin, G.**, Green, M., Kuerer, H., Gong, Y., Browne, D., Hortobagyil, G. N. and Sneige, N. (2007). Comparison of ductal lavage and random periareolar fine needle aspiration as tissue acquisition methods in early breast cancer prevention trials. *Clinical Cancer Research* **13**, 4943–4948.
- 157. Gonzalez, R. J., Buzdar, A. U., Symmans, W. F., Yen, T. W., Broglio, K. R., Lucci, A., Esteva, F. J., **Yin**, **G.** and Kuerer, H. M. (2007). Novel clinical trial designs for treatment of ductal carcinoma in situ of the breast with trastuzumab (herceptin). *Breast Journal* **13**, 72–75.
- 158. Rivera, E., Meyers, C., Groves, M., Valero, V., Francis, D., Arun, B., Broglio, K., **Yin, G.**, Hortobagyi, G. N. and Buchholz, T. (2006). Phase I study of capecitabine in combination with temozolomide in the treatment of patients with brain metastases from breast carcinoma. *Cancer* **107**, 1348–1354.
- 159. Sneige, N., Liu, B., **Yin, G.** and Arun, B. K. (2006). Correlation of cytologic findings and chromosomal instability detected by fluorescence in situ hybridization in breast fine-needle aspiration specimens from women at high risk for breast cancer. *Modern Pathology* **19**, 622–629.
- 160. Klos, K. S., Sun, M., Tan, M., Zhou, X., Li, P., Yang, W., Yin, G. and Yu, D. (2006). ErbB2 increases VEGF protein synthesis via activation of the mTOR/p70S6K pathway leading to increased angiogenesis and spontaneous metastasis of human breast cancer cells. *Cancer Research* 66, 2028–2037.
- 161. Tan, M., Li, P., Sun, M., Yin, G. and Yu, D. (2006). Upregulation and activation of PKC α by ErbB2 through Src promotes breast cancer cell invasion that can be blocked by combined treatment with PKC α and Src inhibitors. *Oncogene* 25, 3286–3295.
- 162. Hanrahan, E. O., Broglio, K. R., Buzdar, A. U., Theriault, R. L., Valero, V., Cristofanilli, M., Yin, G., Kau, S.-W., Hortobagyi, G. N. and Rivera, E. (2005). Combined-modality treatment for isolated recurrences of breast carcinoma. *Cancer* 104, 1158–1171.
- 163. Caplan, D., Cai, J., Yin, G. and White, A. (2005). Root canal filled versus non-root canal filled teeth: A retrospective comparison of survival times. *Journal of Public Health Dentistry* 65, 90–96.