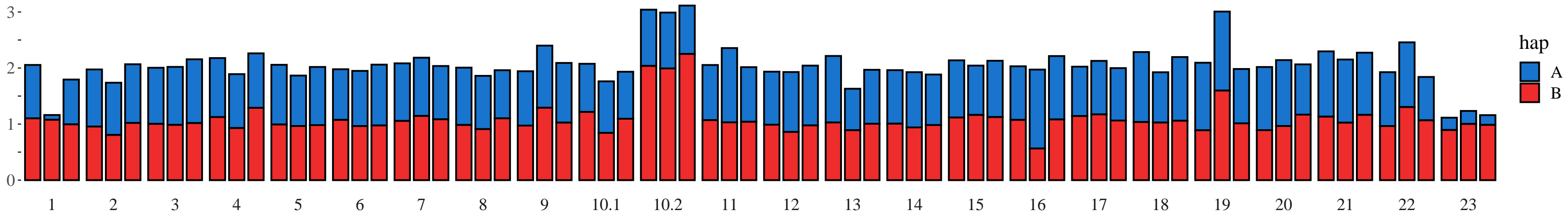
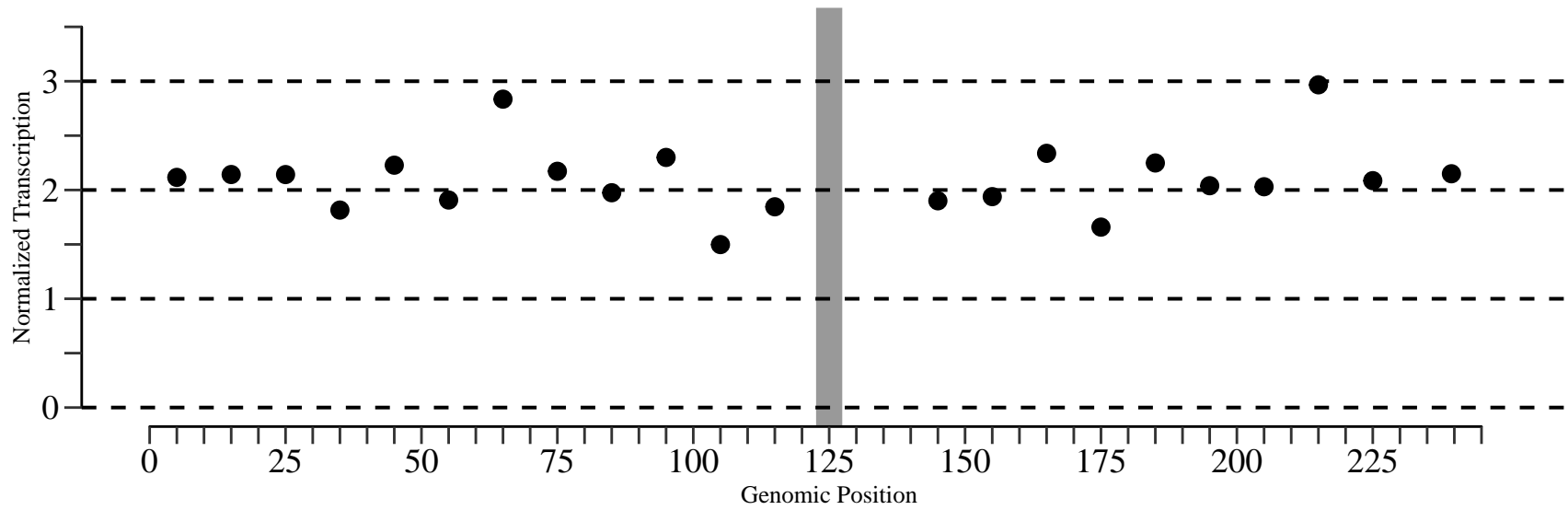


	generation	family_ids	A_id	B_id	chr	A_TPM	B_TPM	c1_TPM	c2_TPM	MN_hap
109	Gen2	F227	F227.3	F227.2	1	1.794819	1.162502	2.055469	NA	A
110	Gen2	F227	F227.3	F227.2	11	2.015567	2.356191	2.054174	NA	A
111	Gen2	F227	F227.3	F227.2	16	2.213689	1.972506	2.031196	NA	NA
112	Gen2	F227	F227.3	F227.2	19	1.983166	3.007391	2.094471	NA	NA

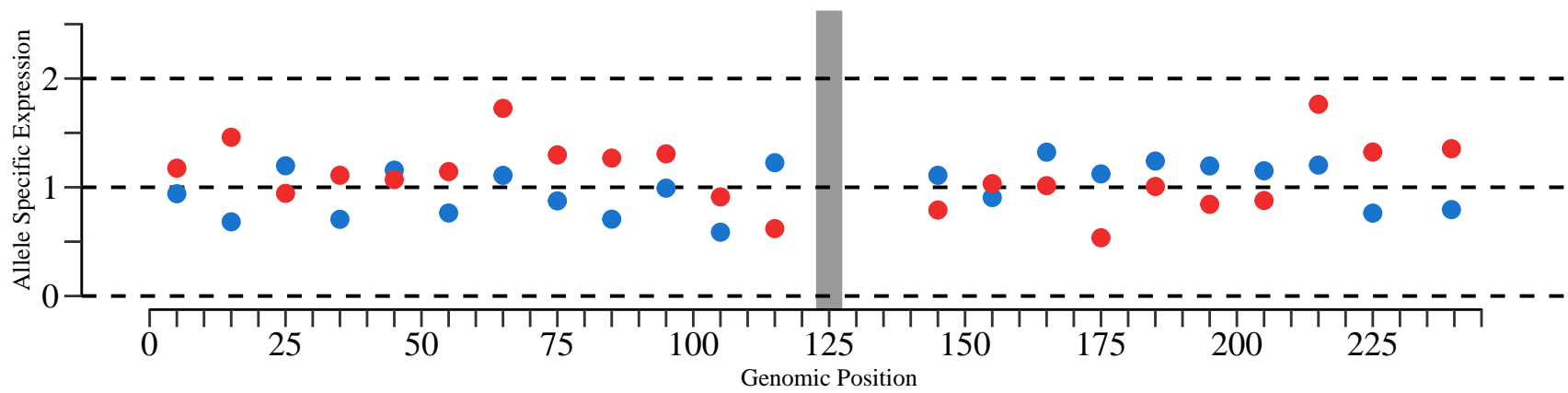
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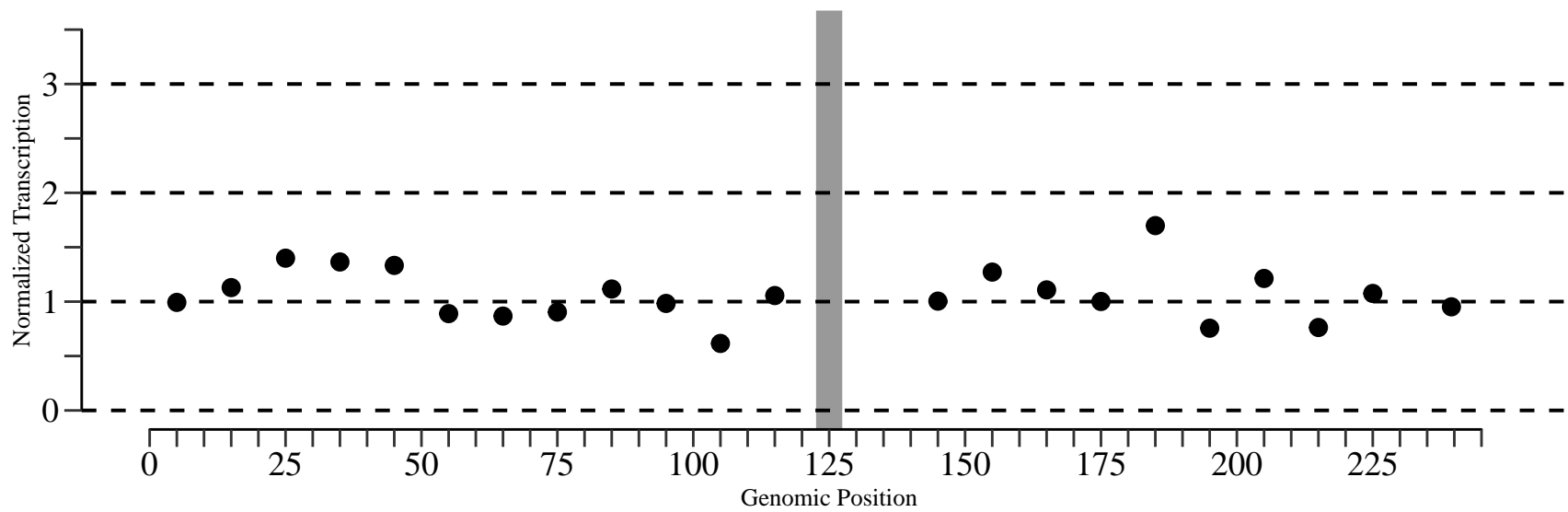
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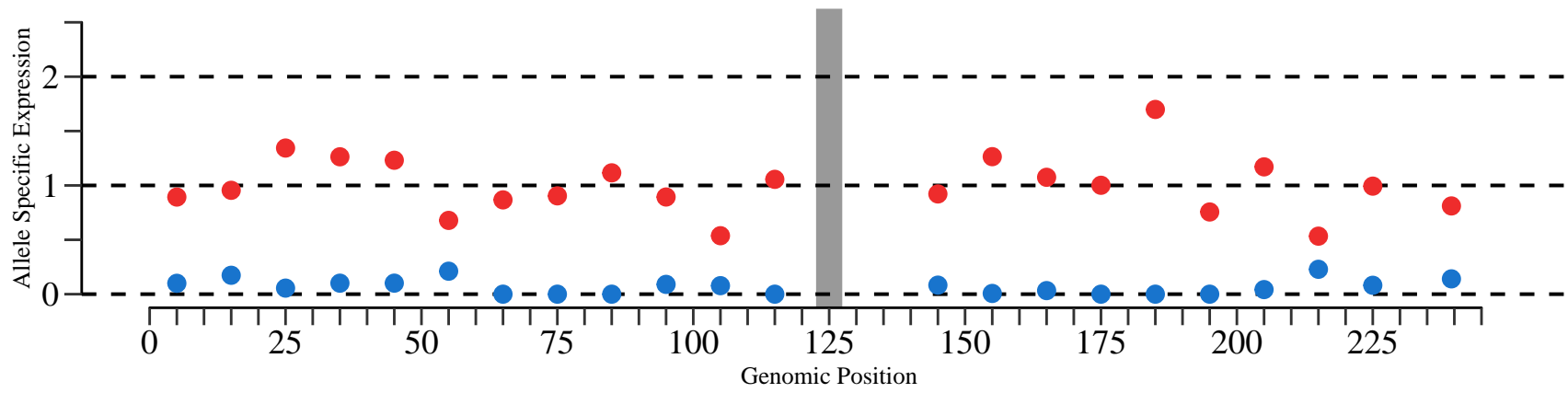
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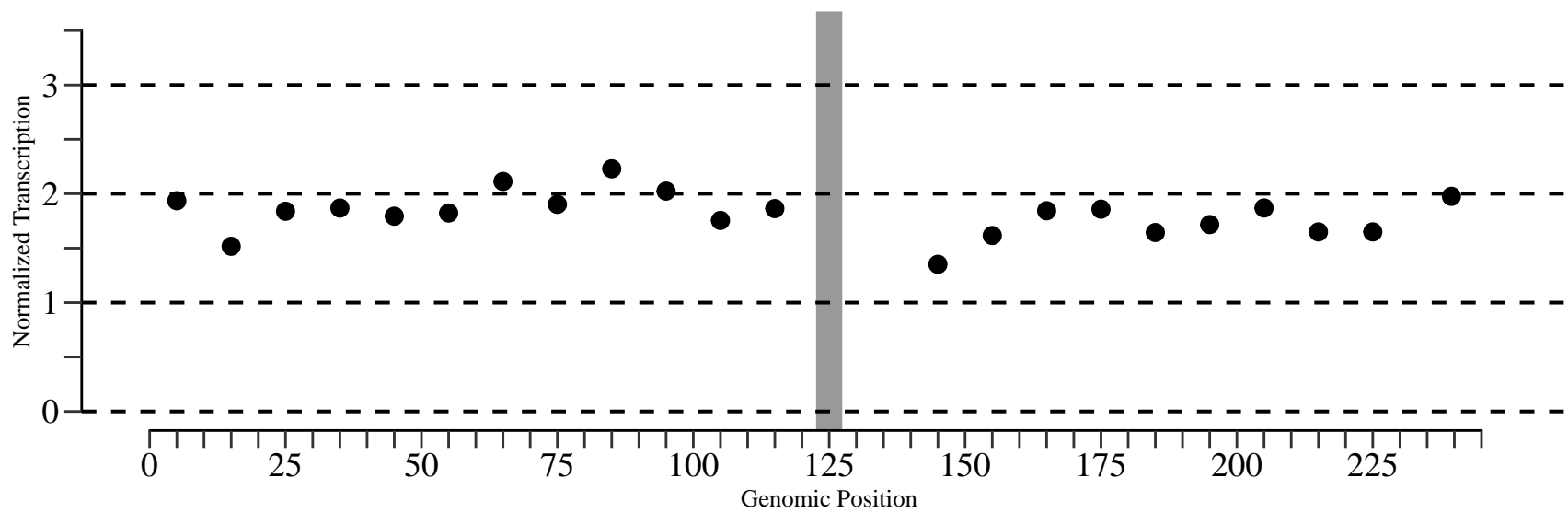
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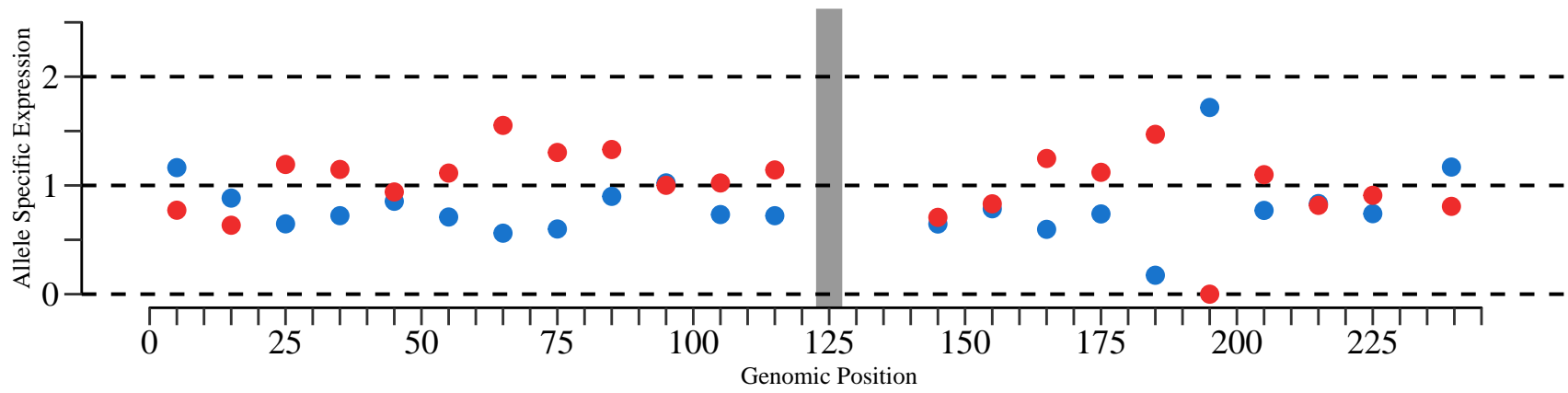
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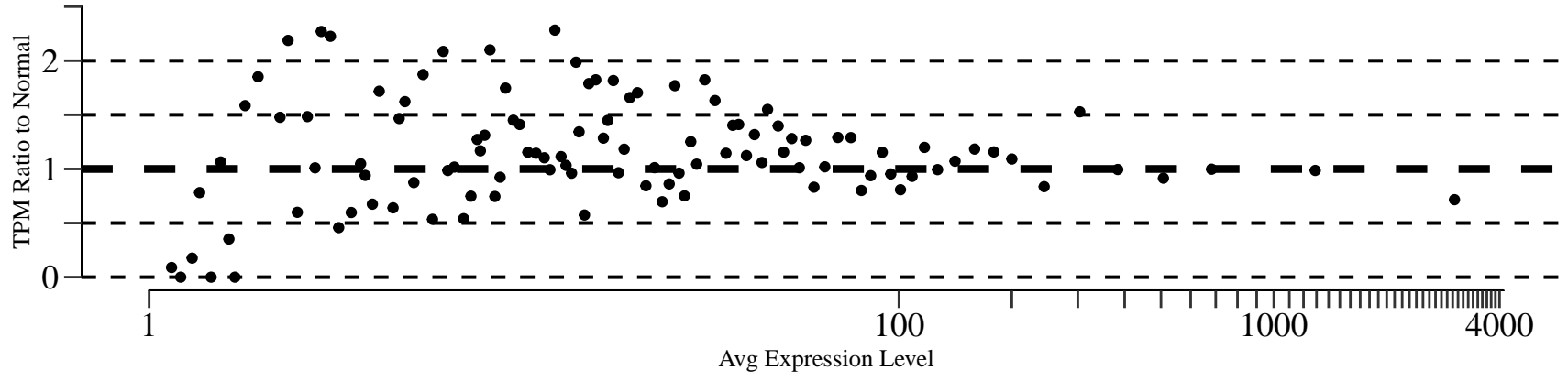
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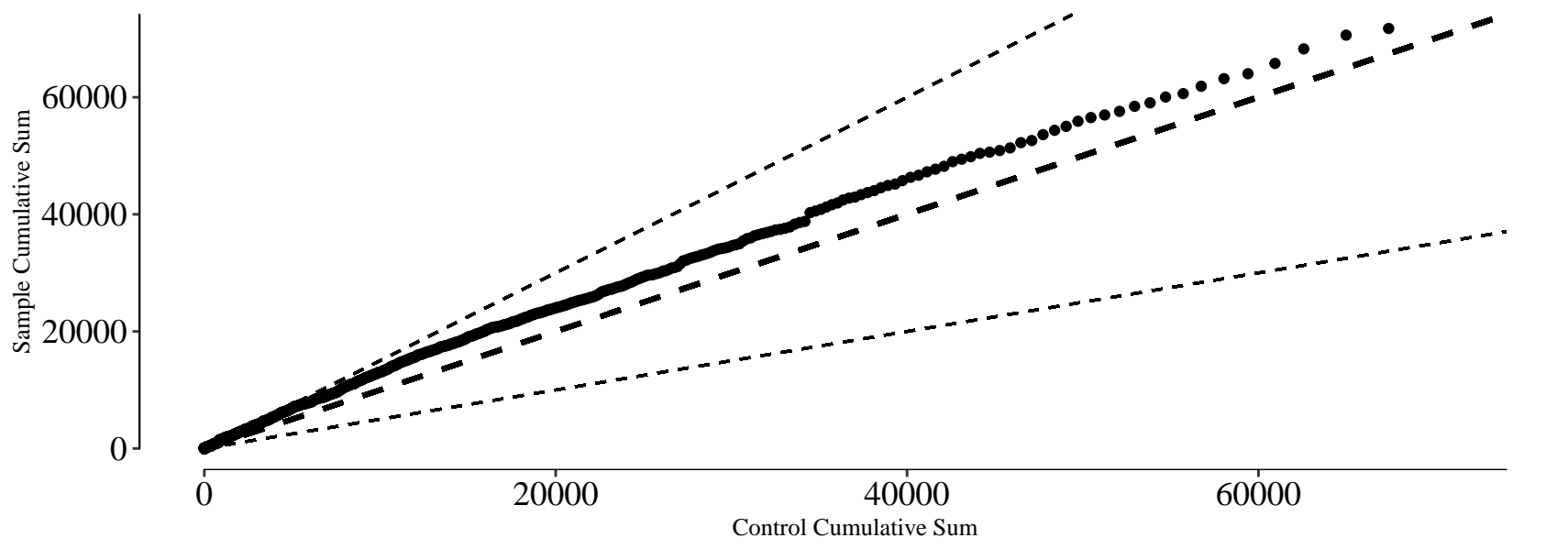
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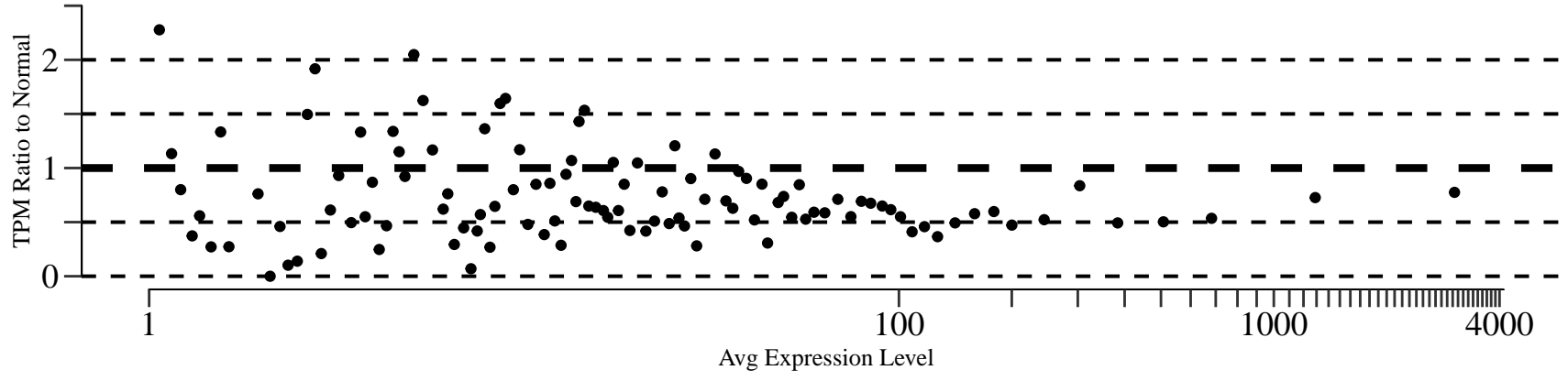
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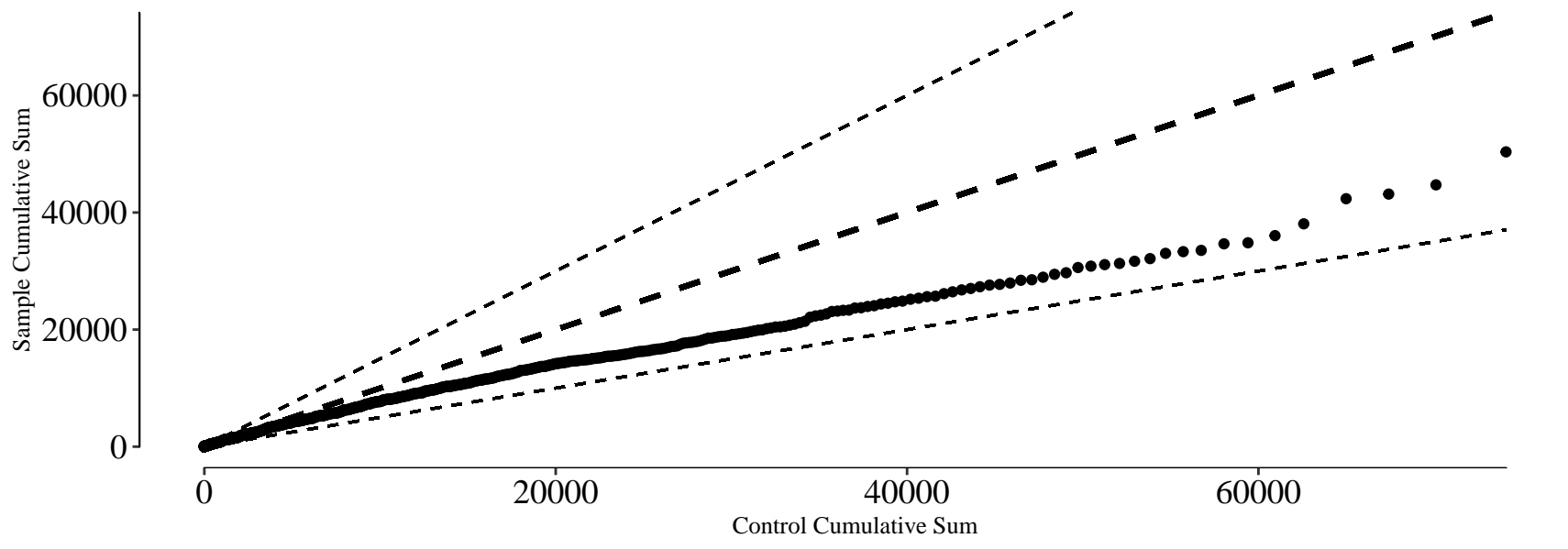
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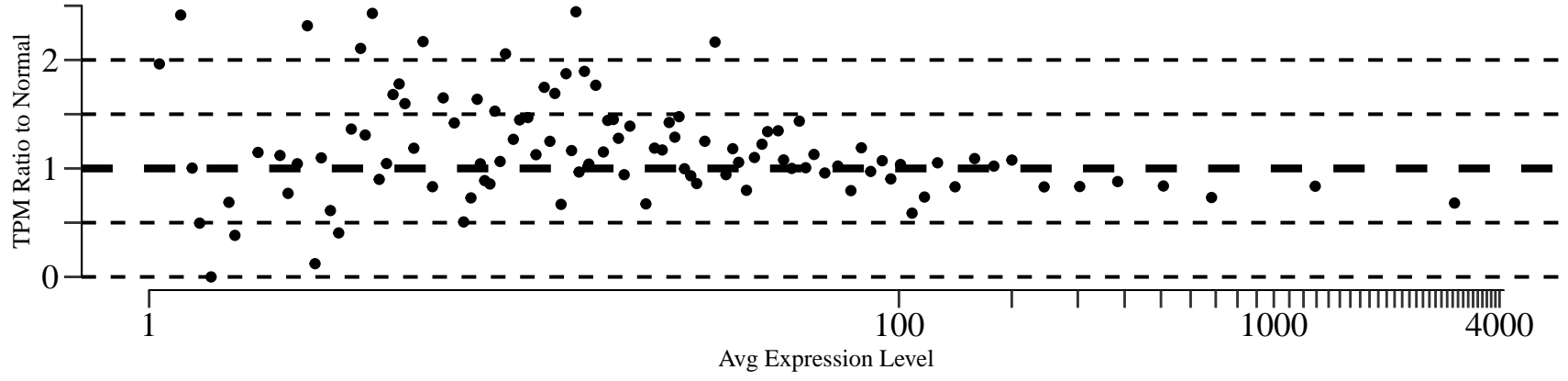
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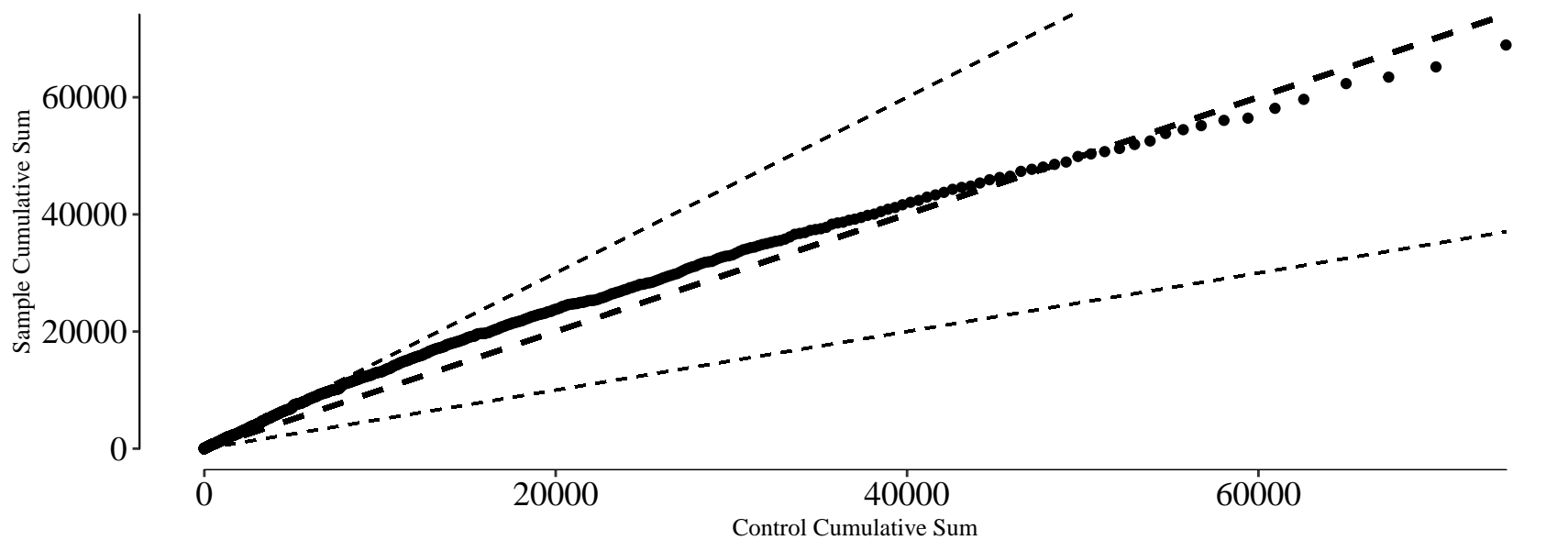
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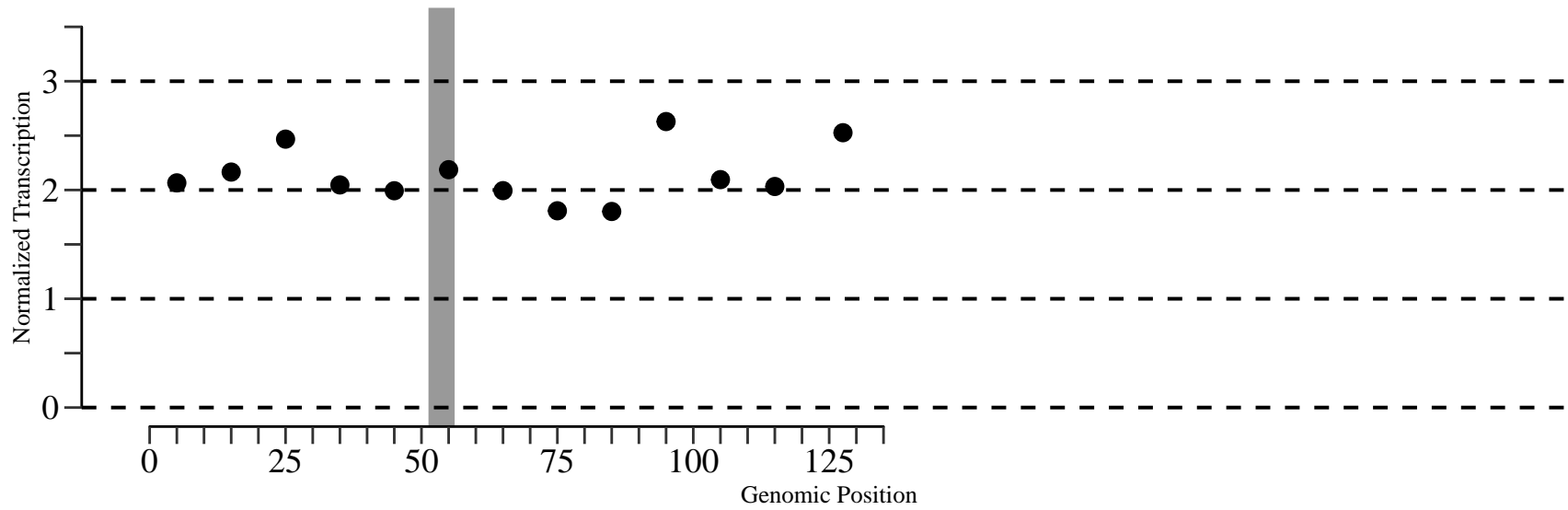
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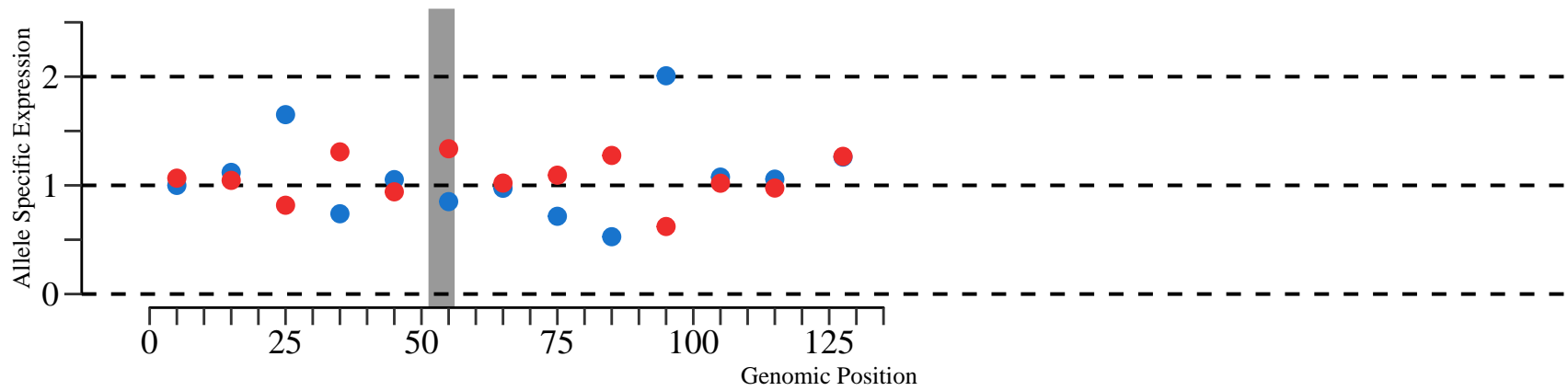
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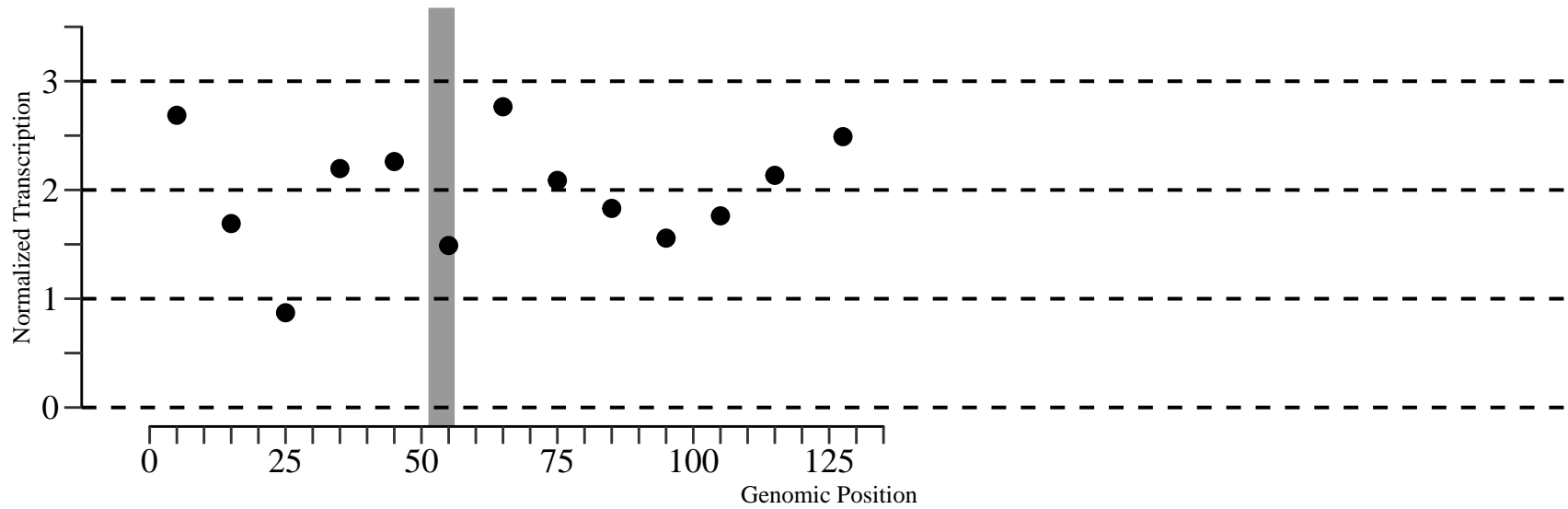
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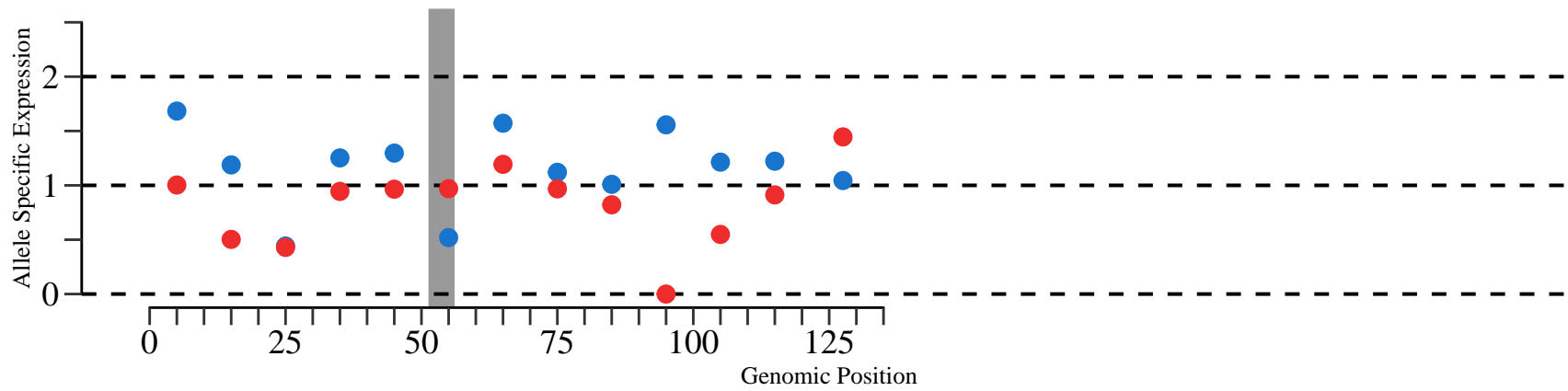
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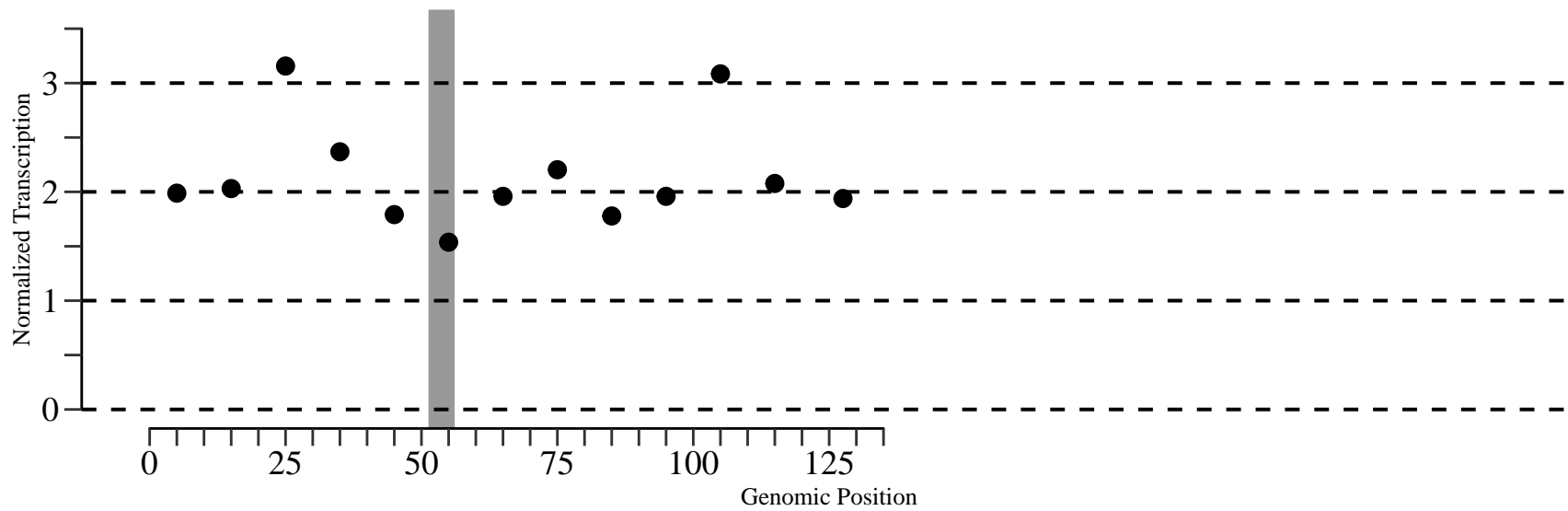
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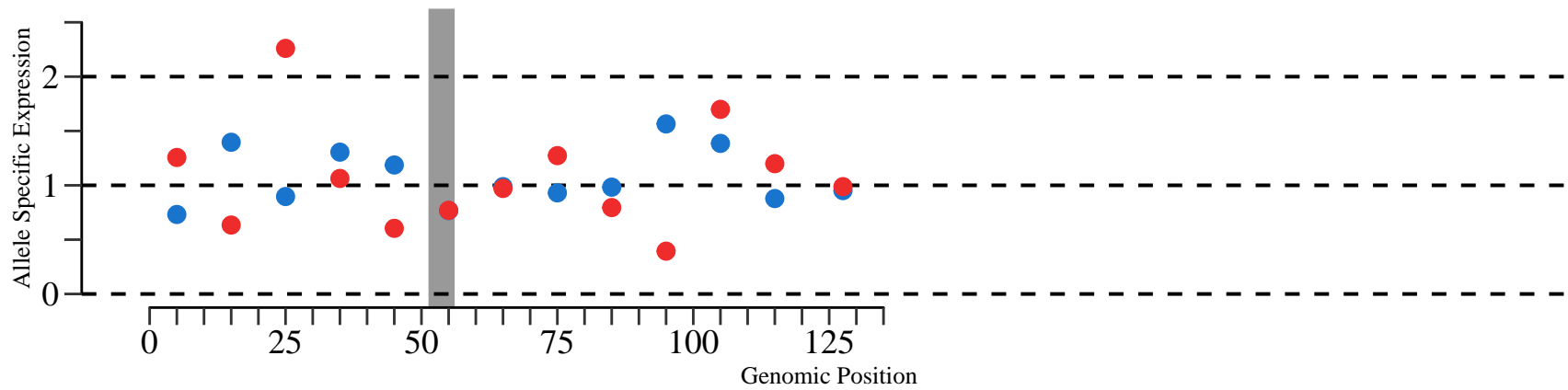
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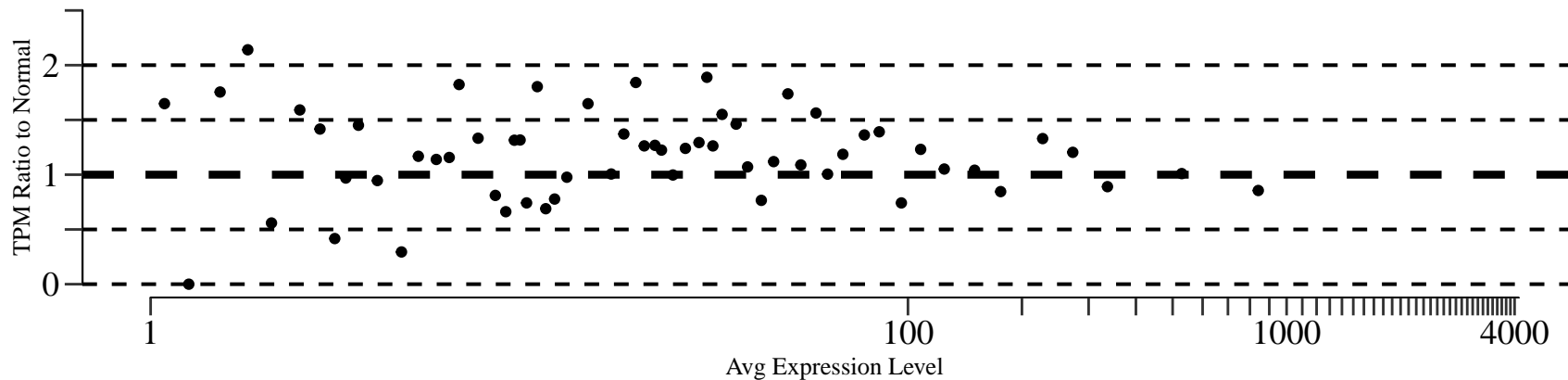
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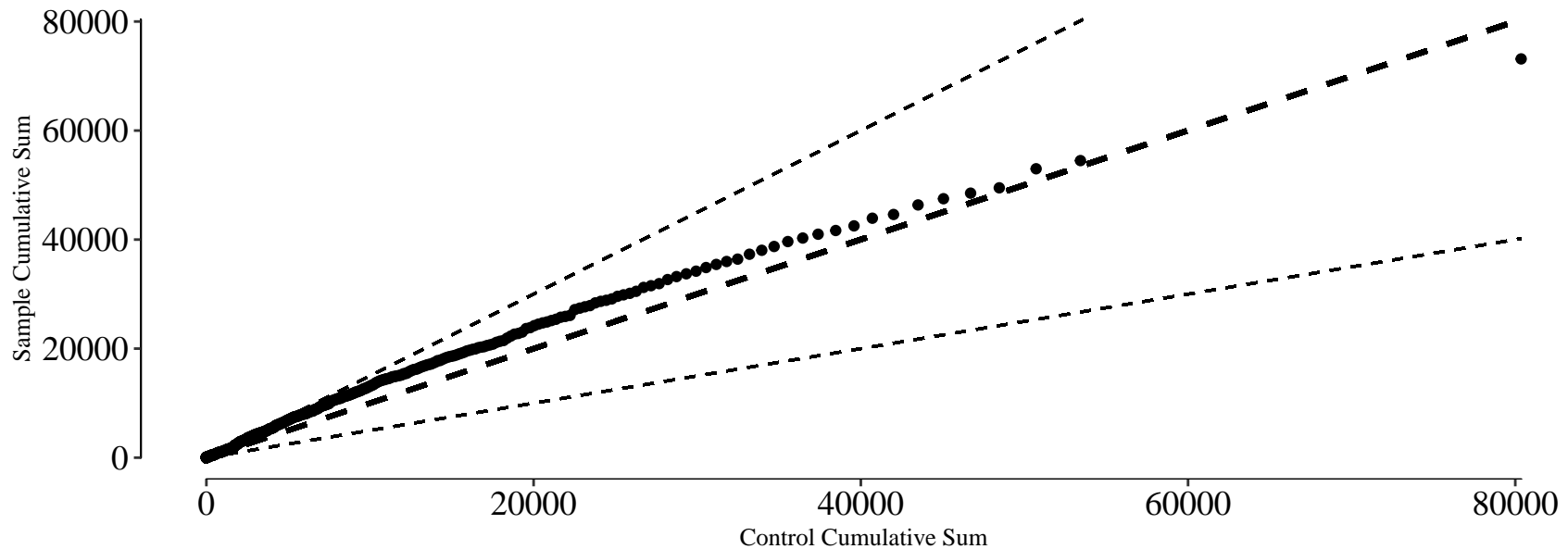
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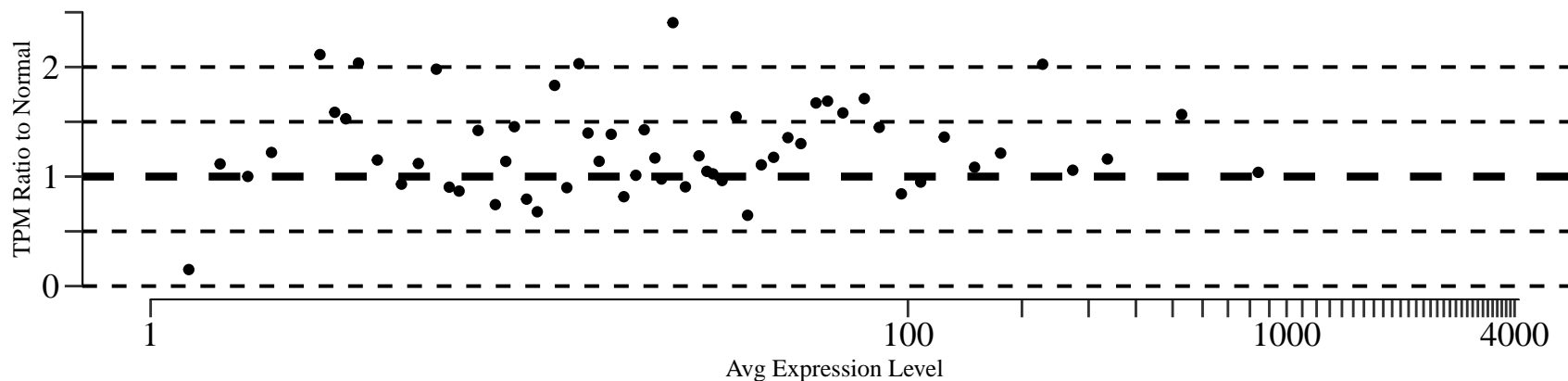
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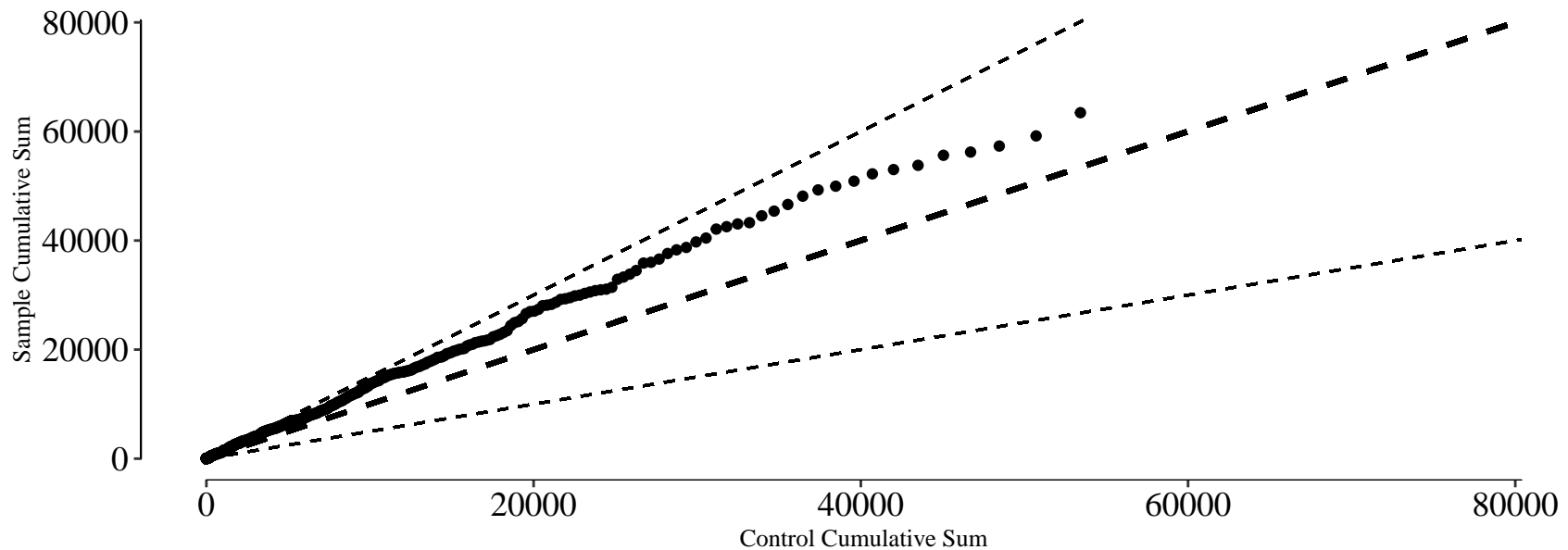
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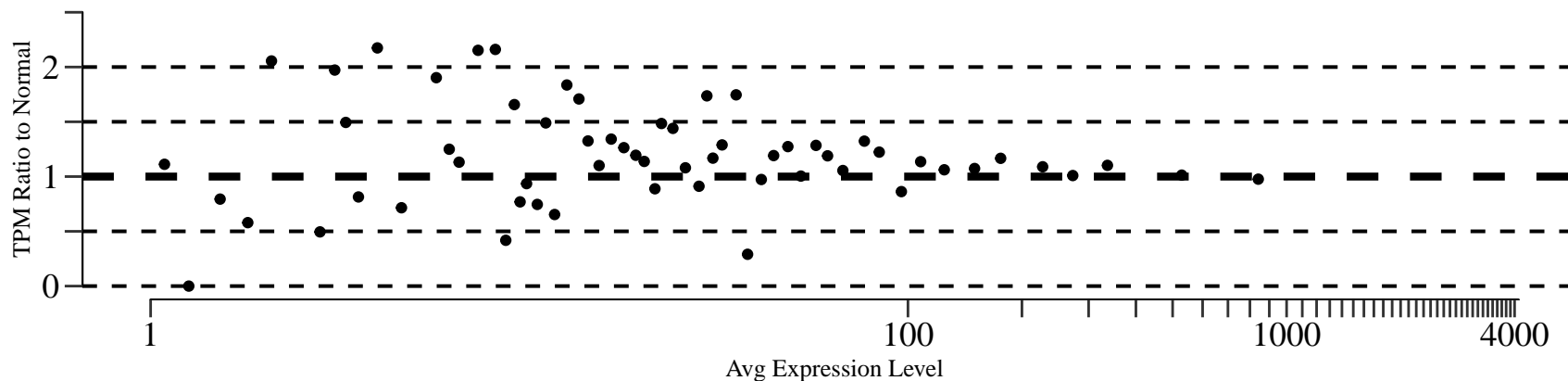
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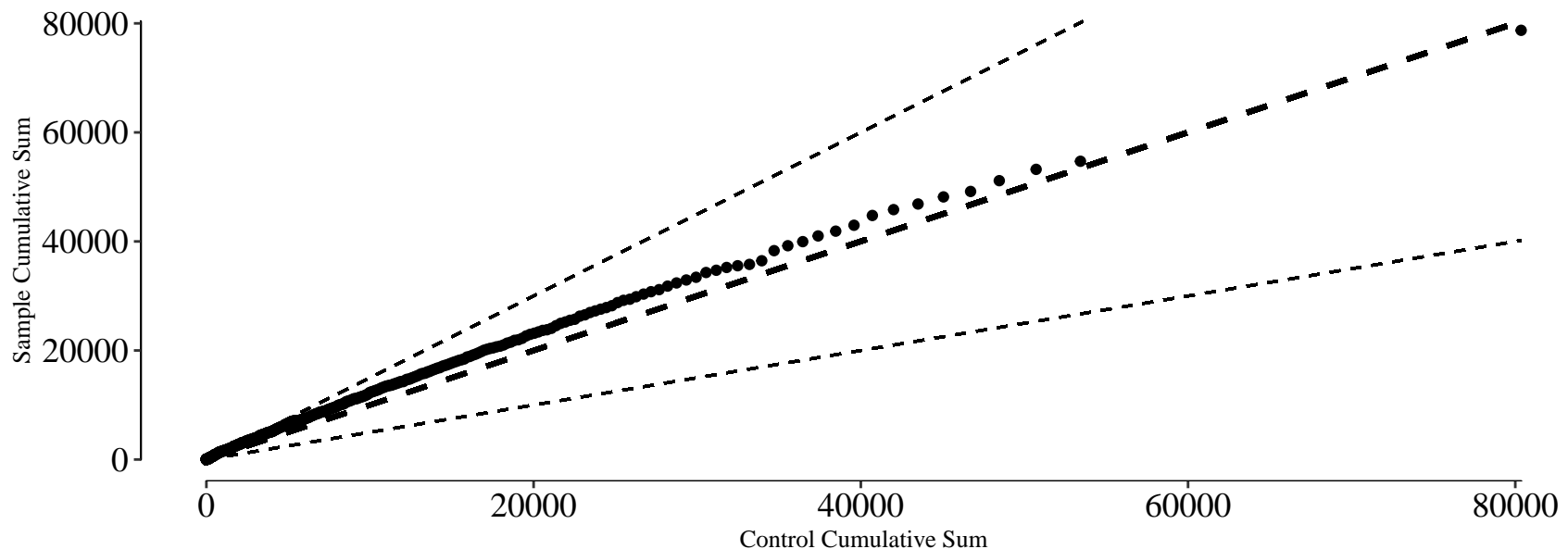
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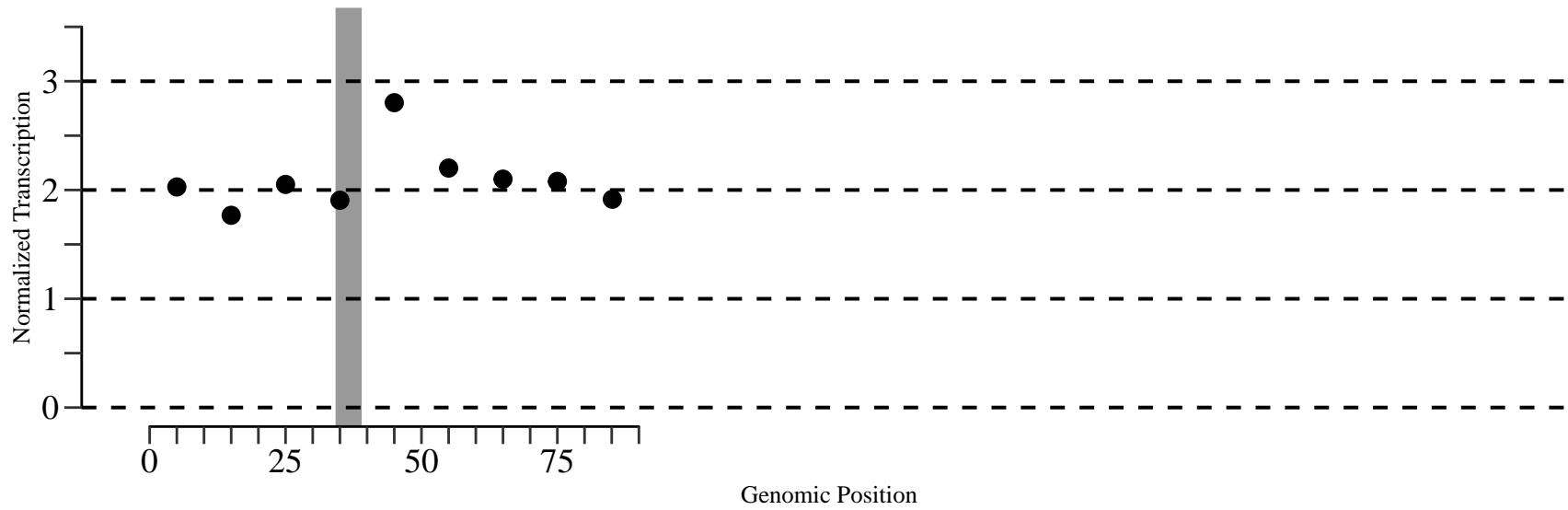
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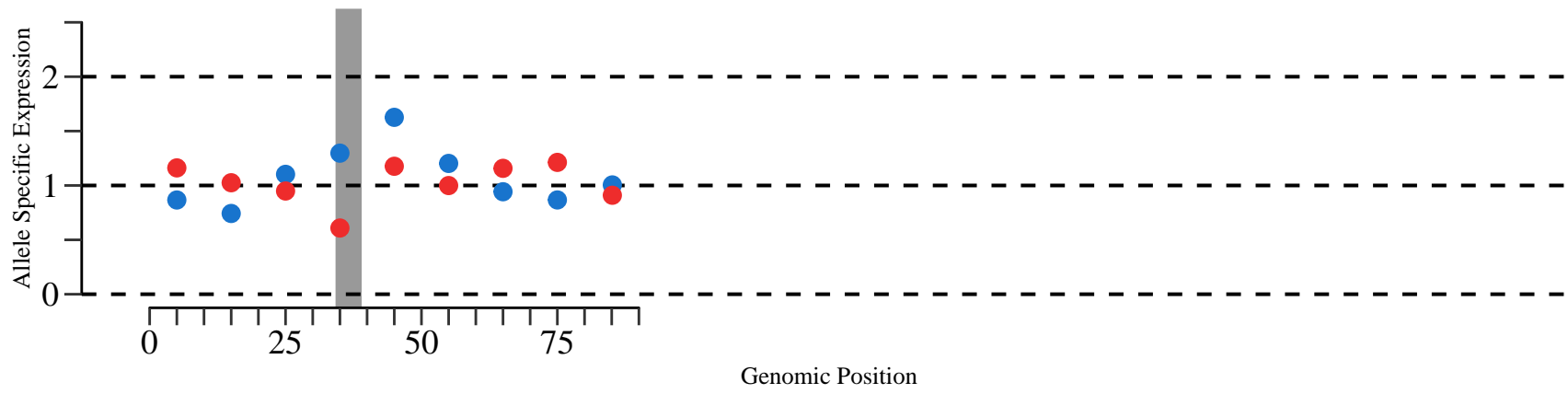
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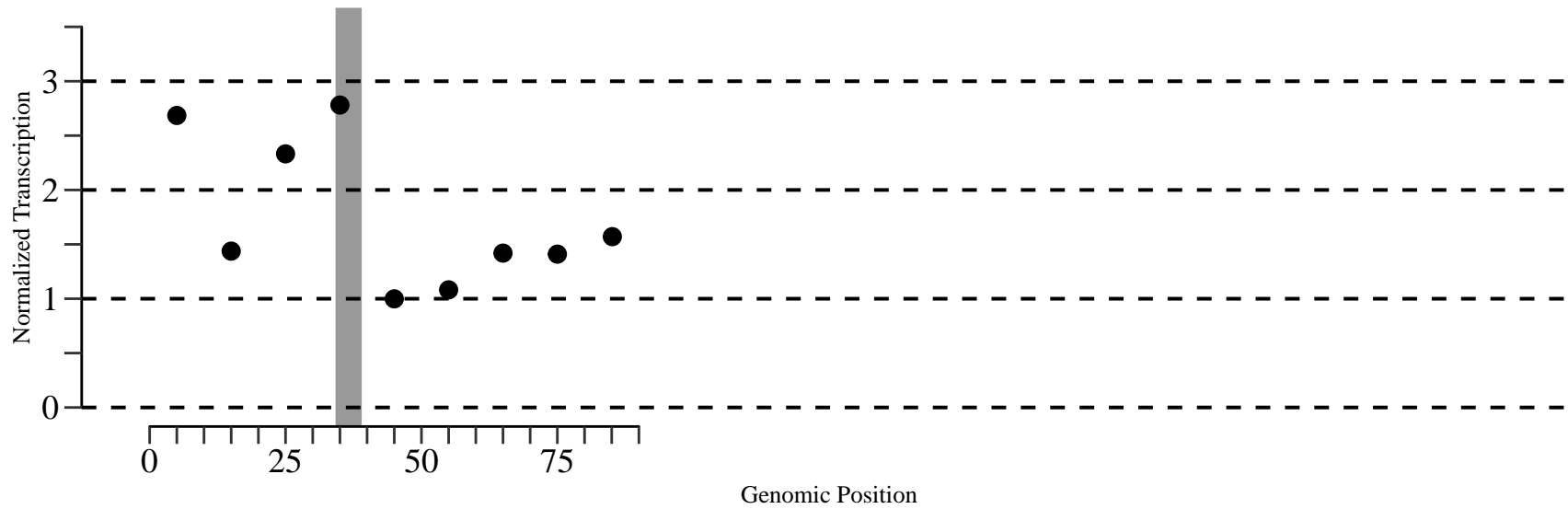
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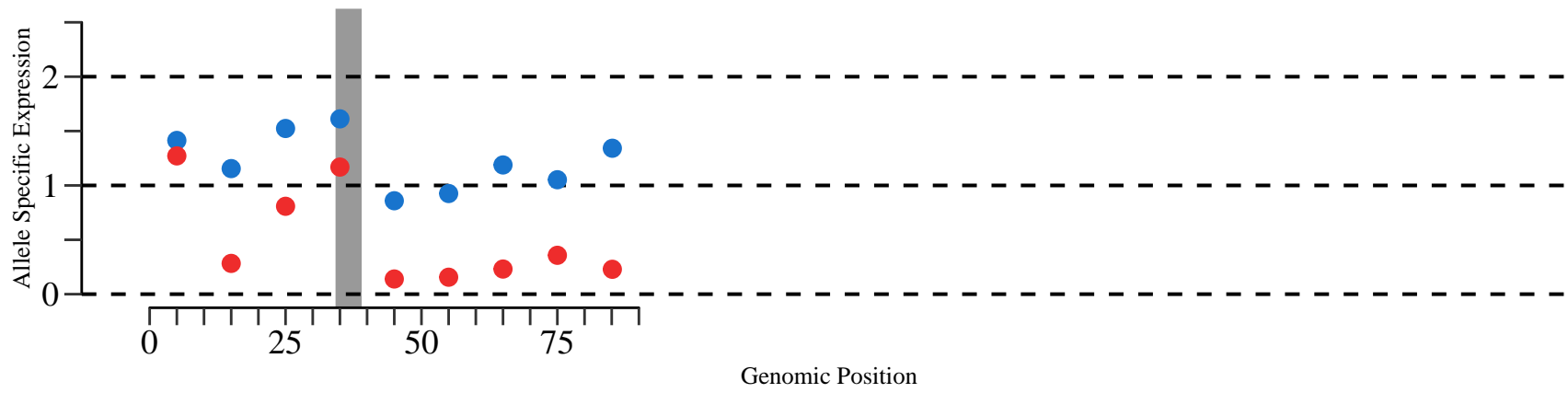
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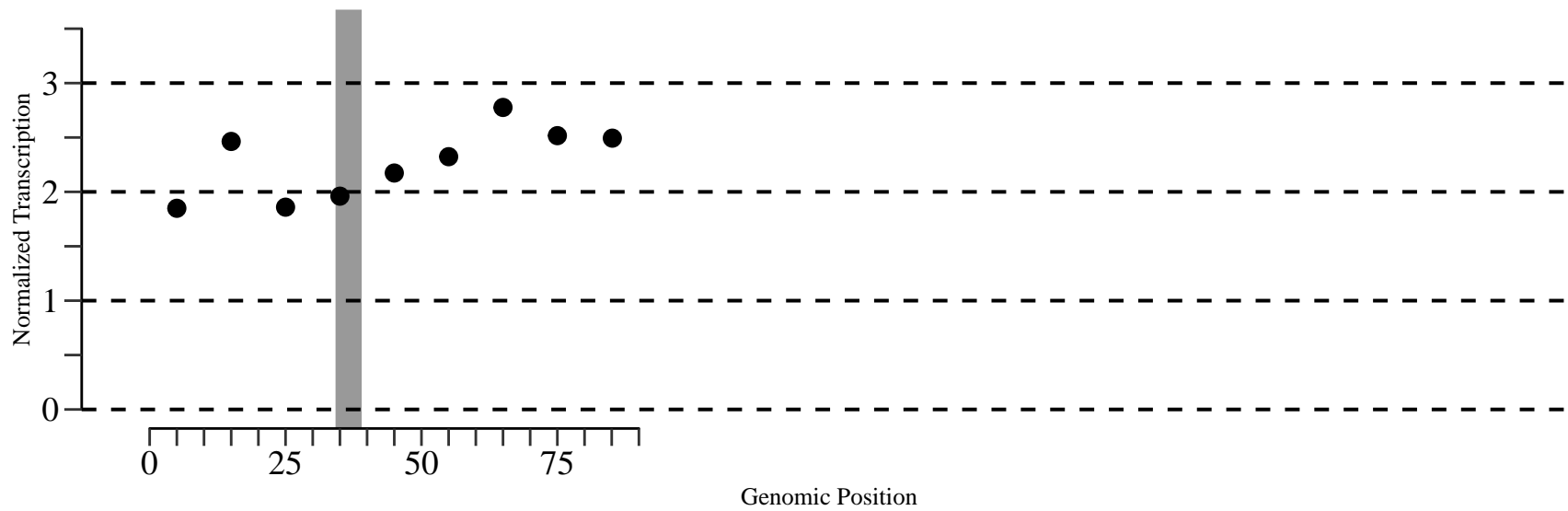
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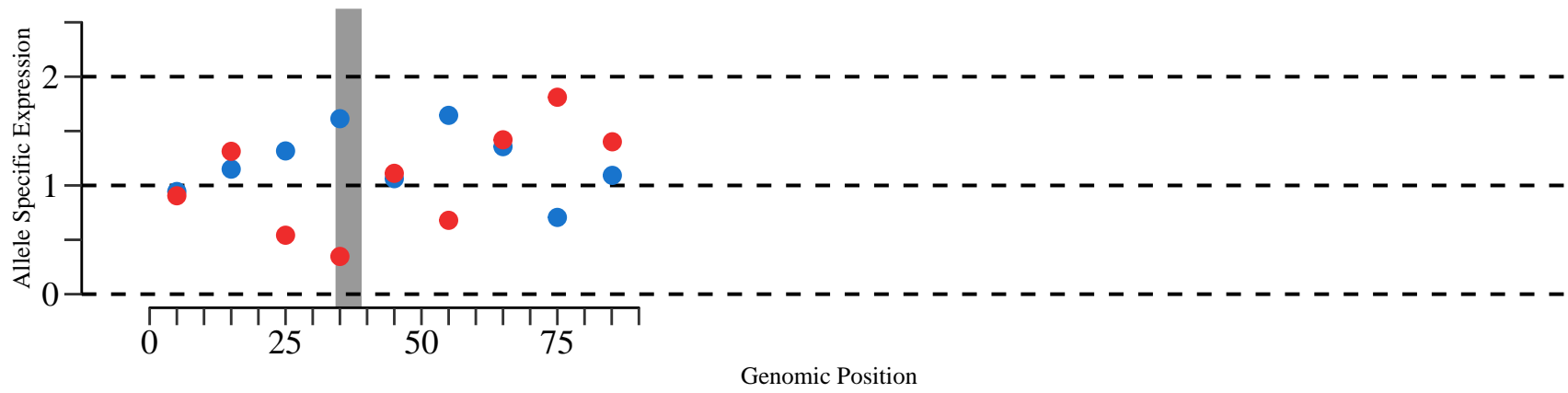
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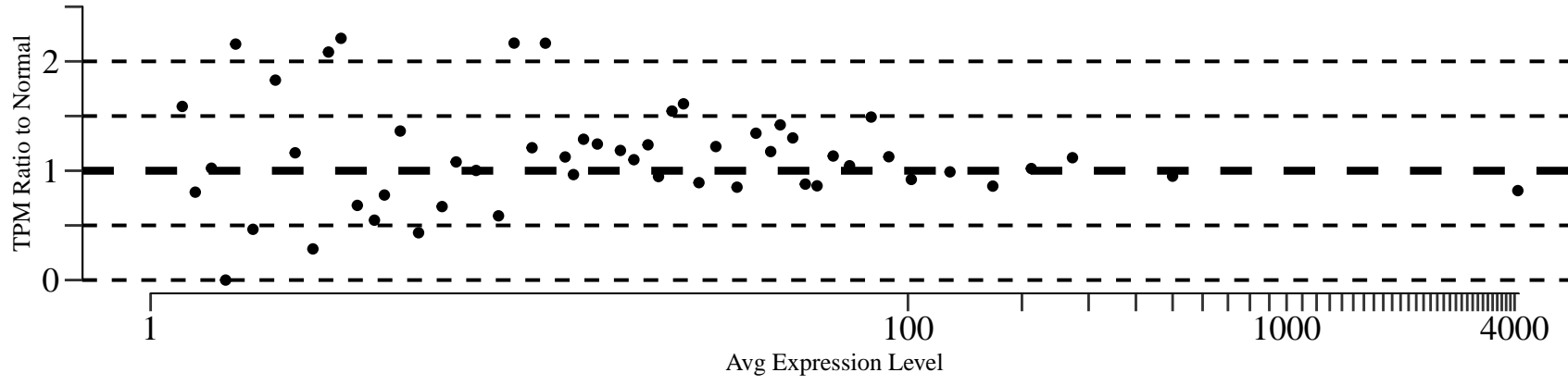
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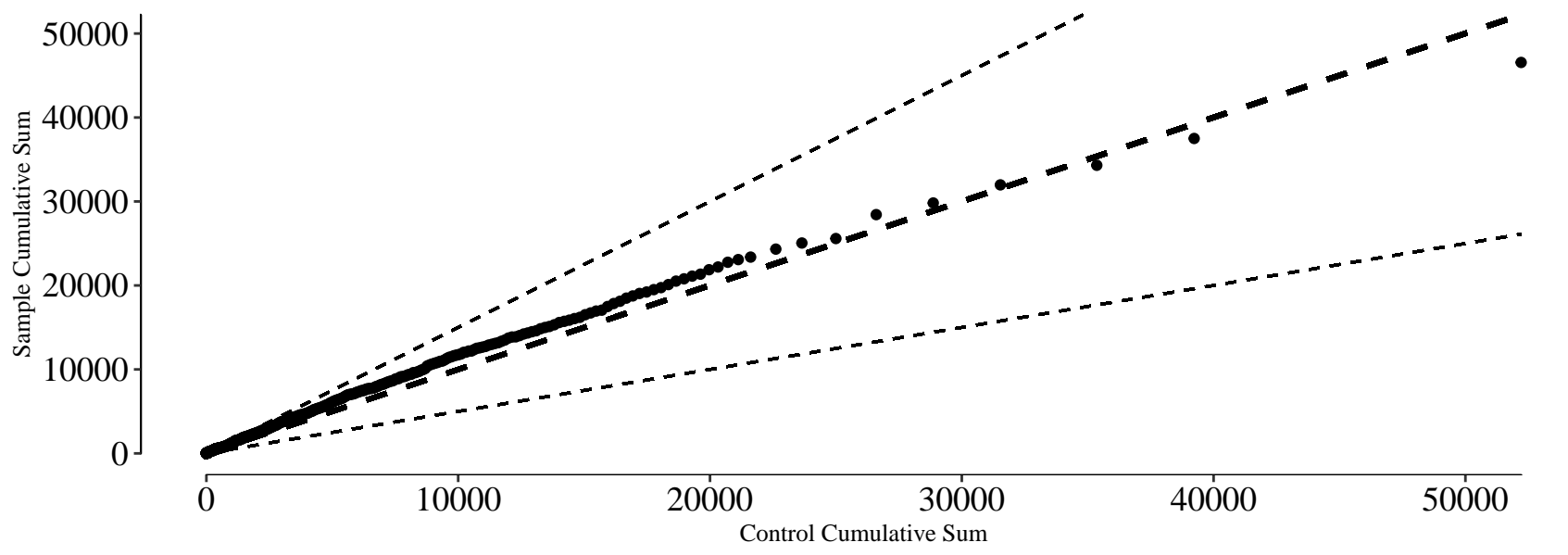
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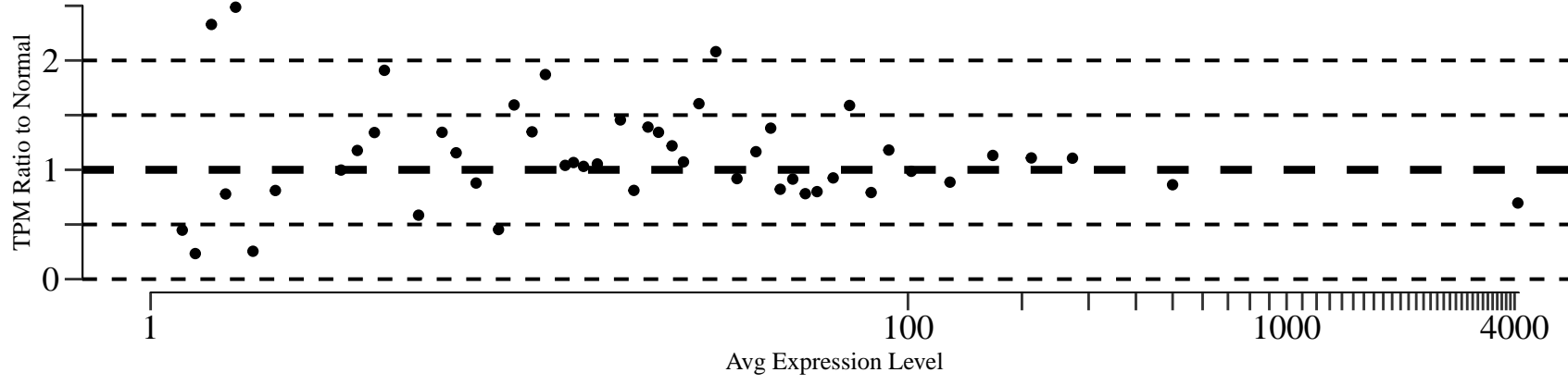
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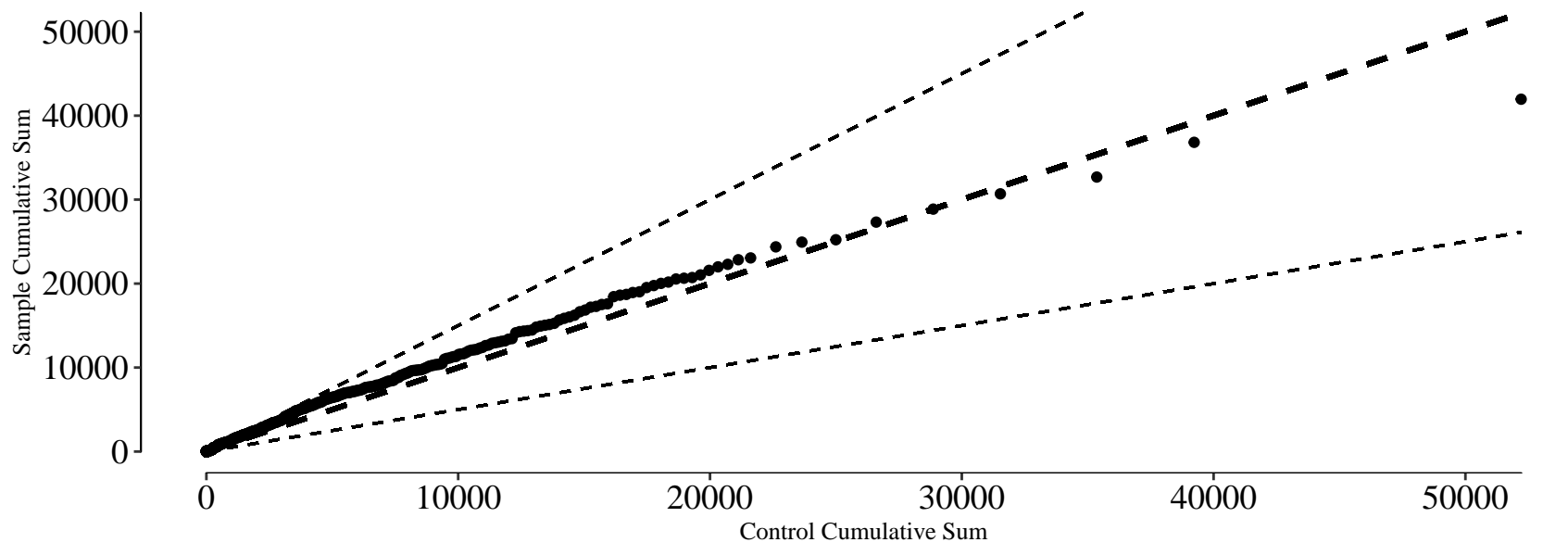
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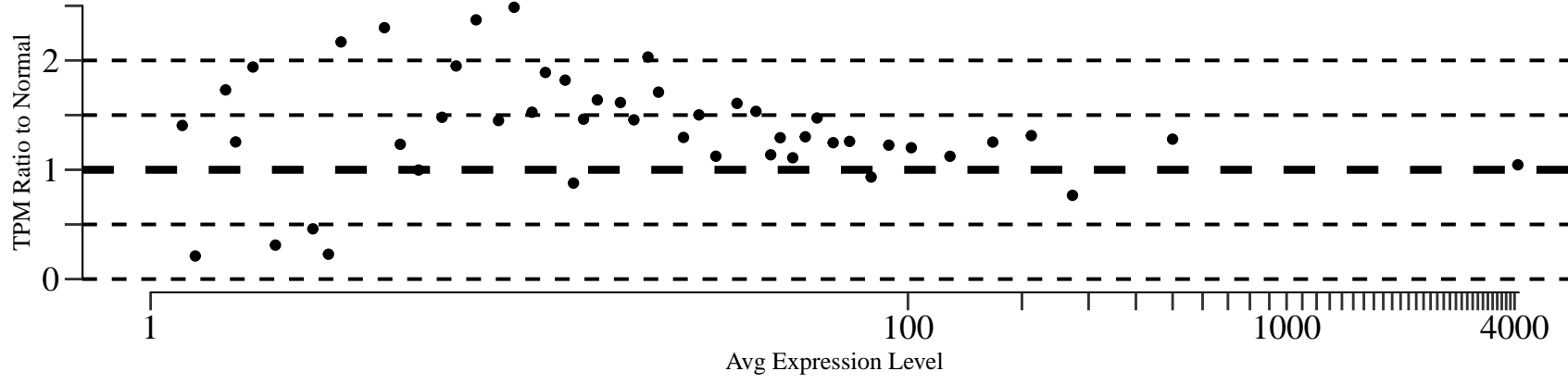
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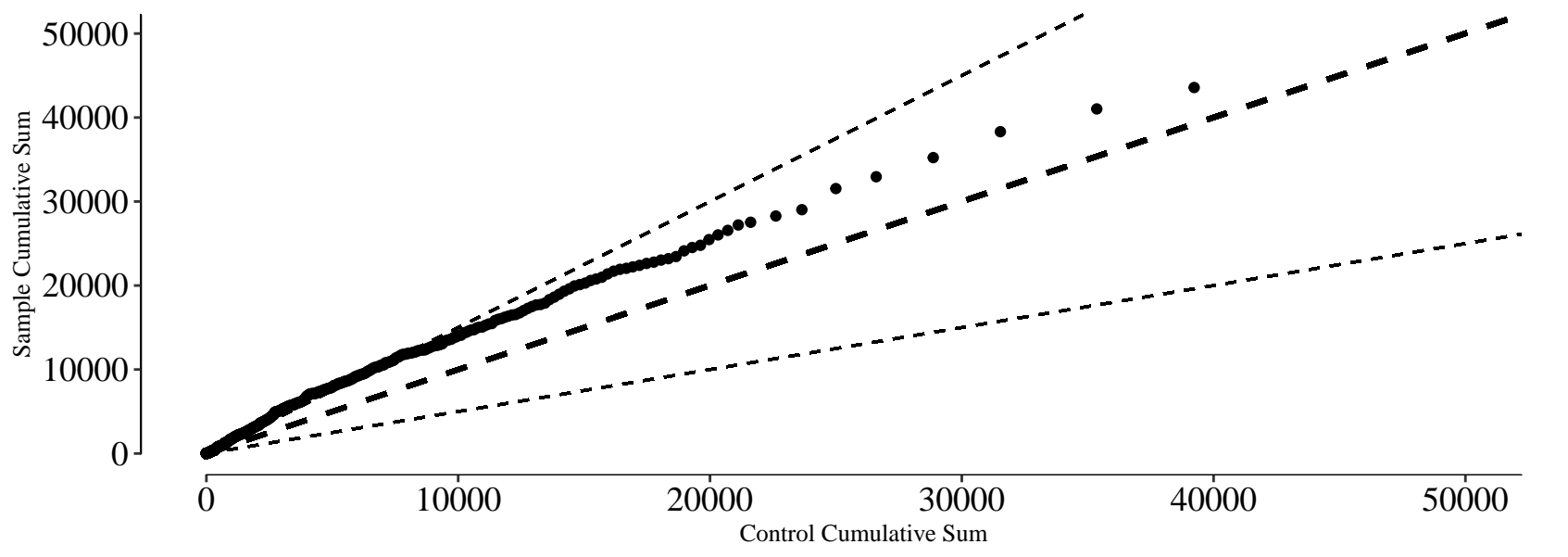
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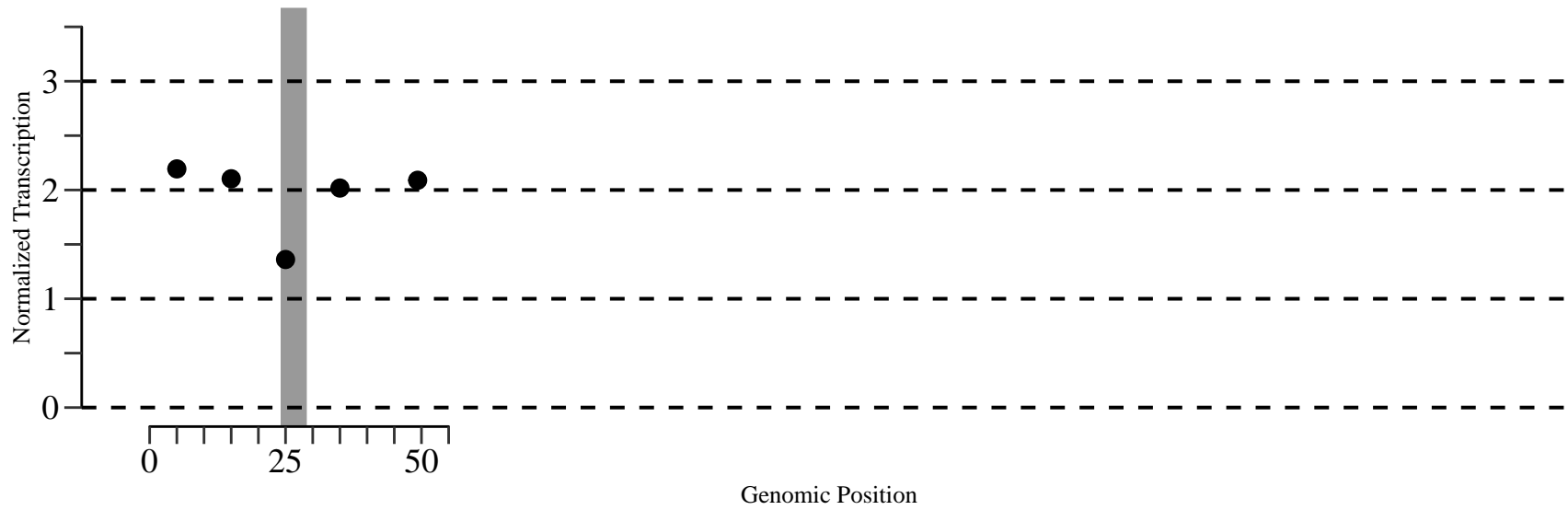
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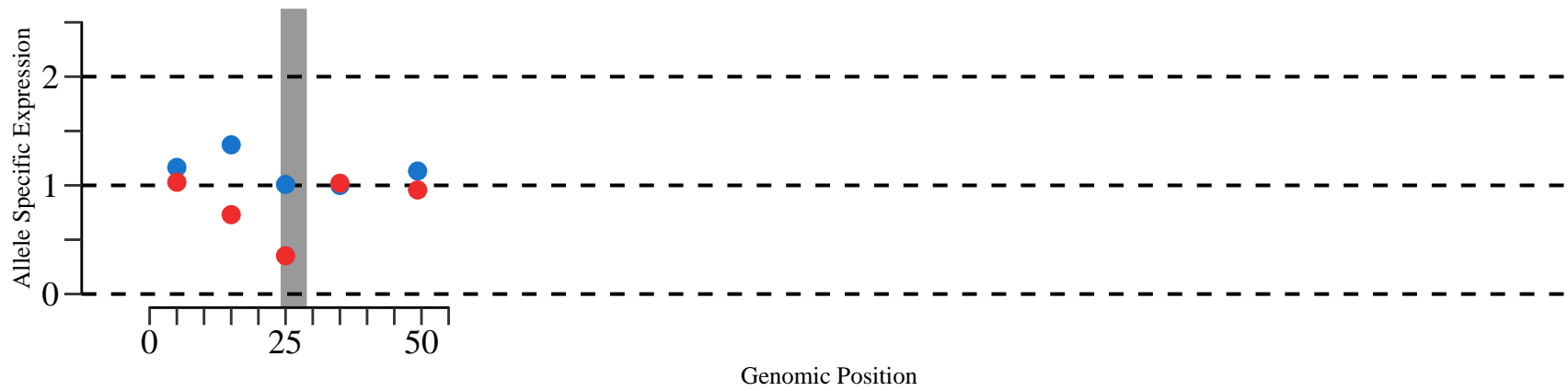
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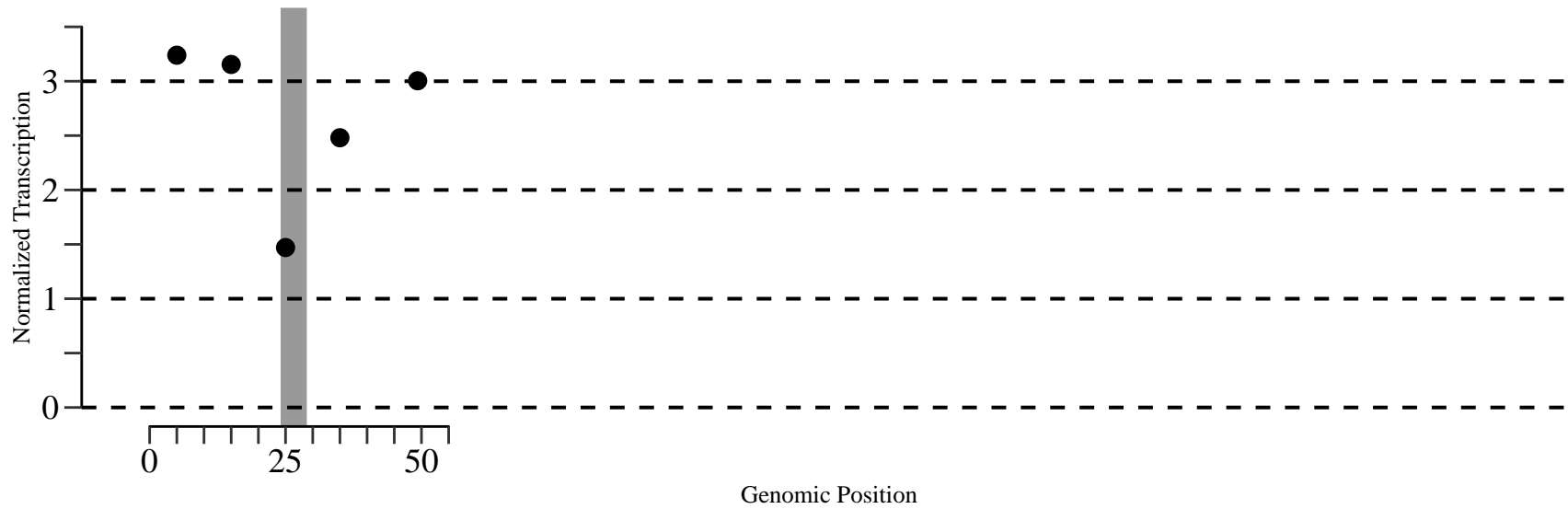
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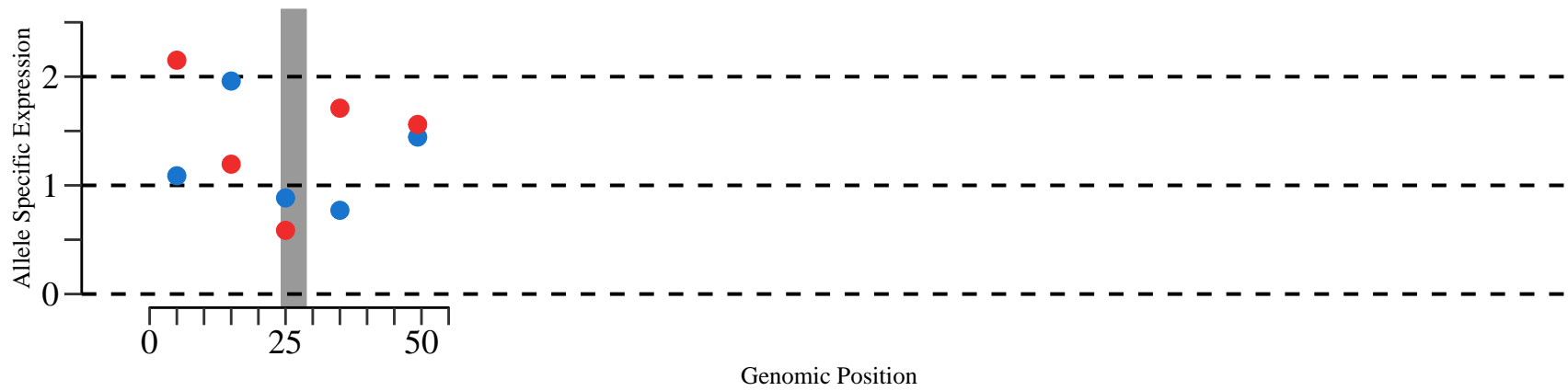
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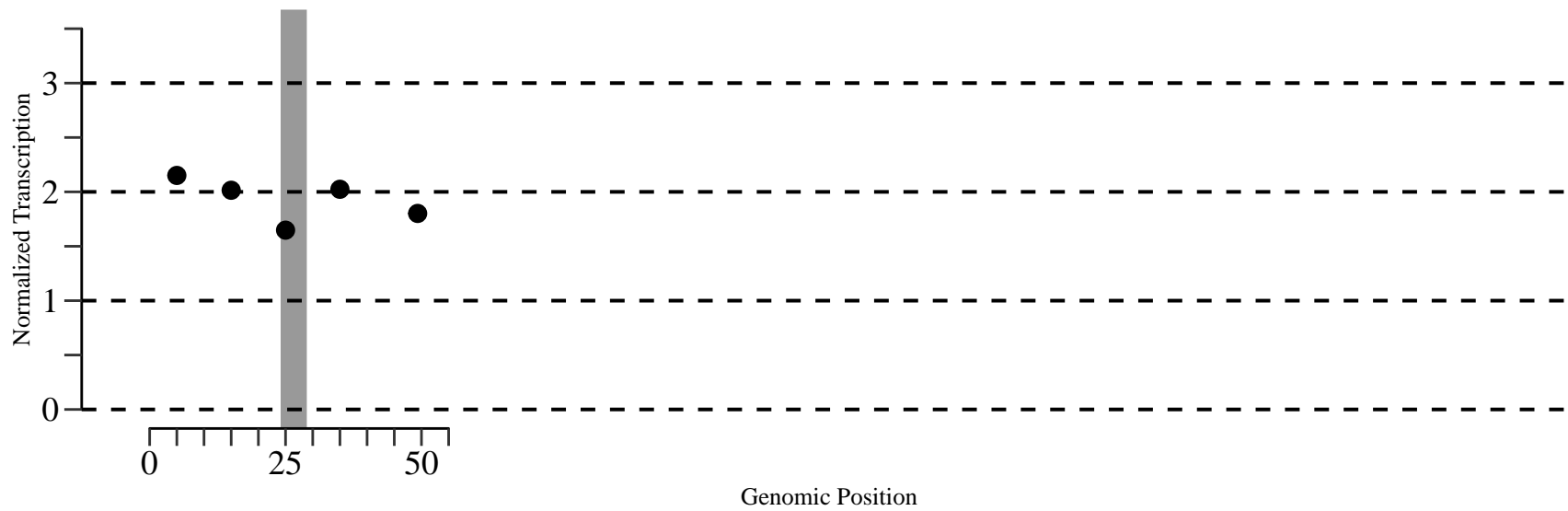
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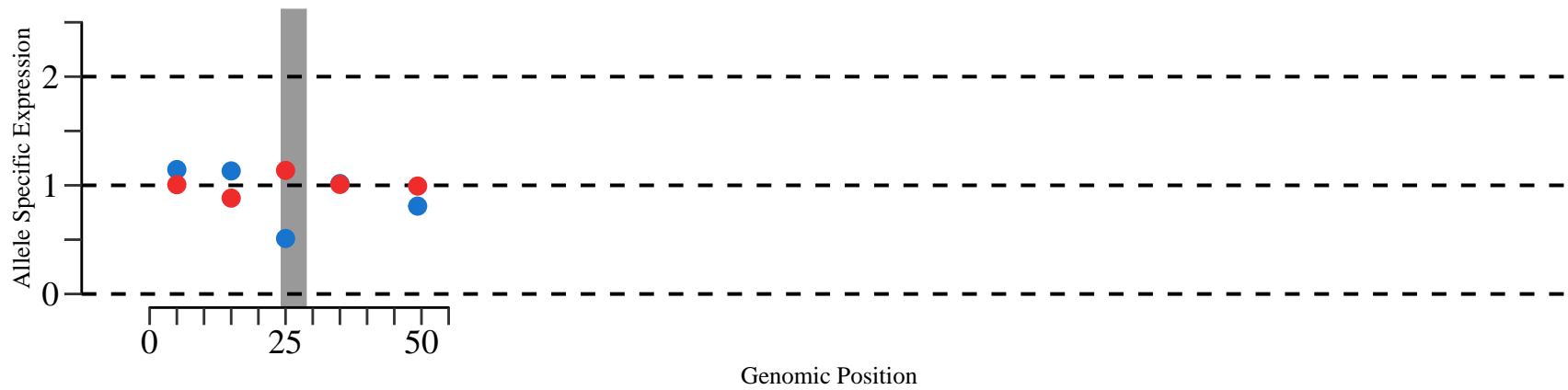
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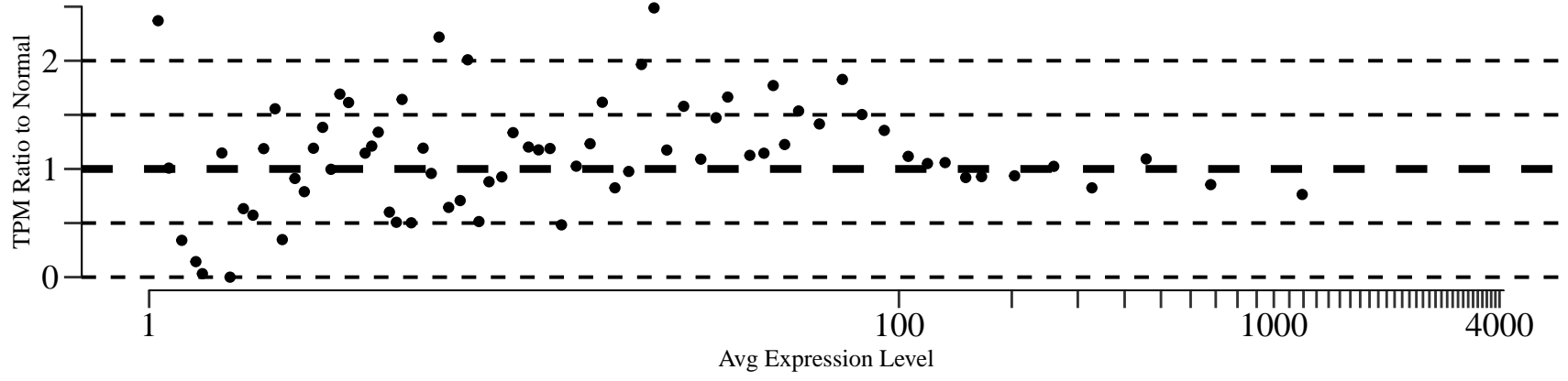
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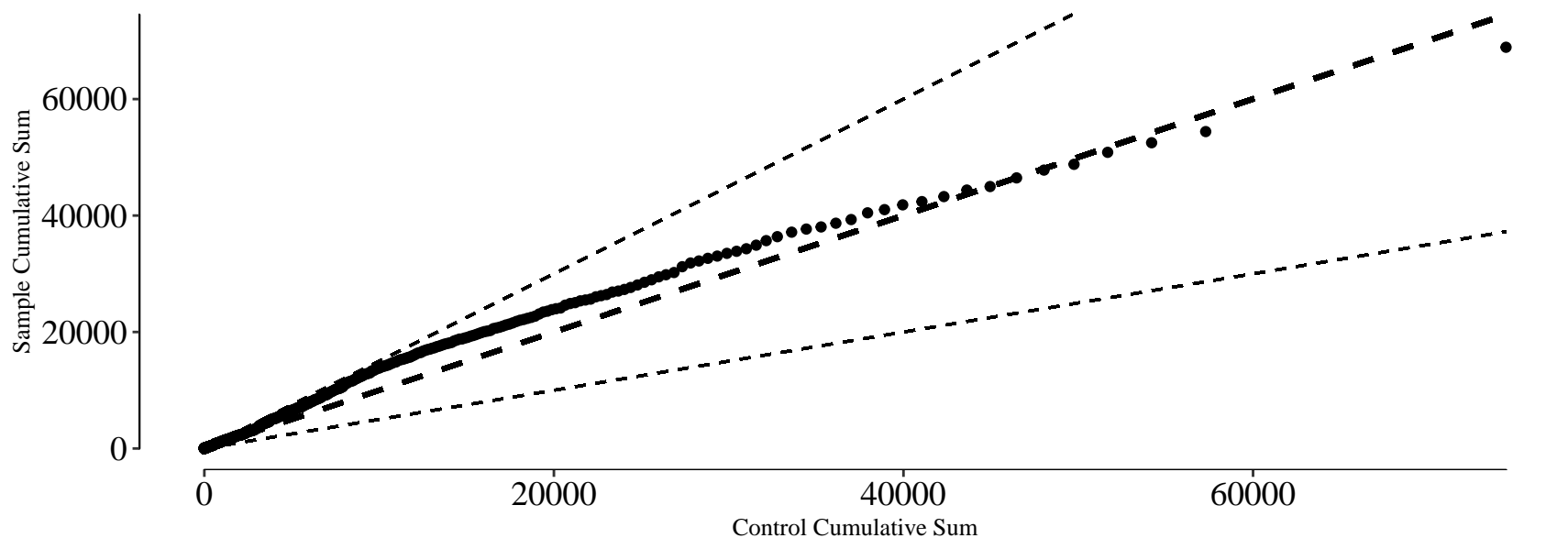
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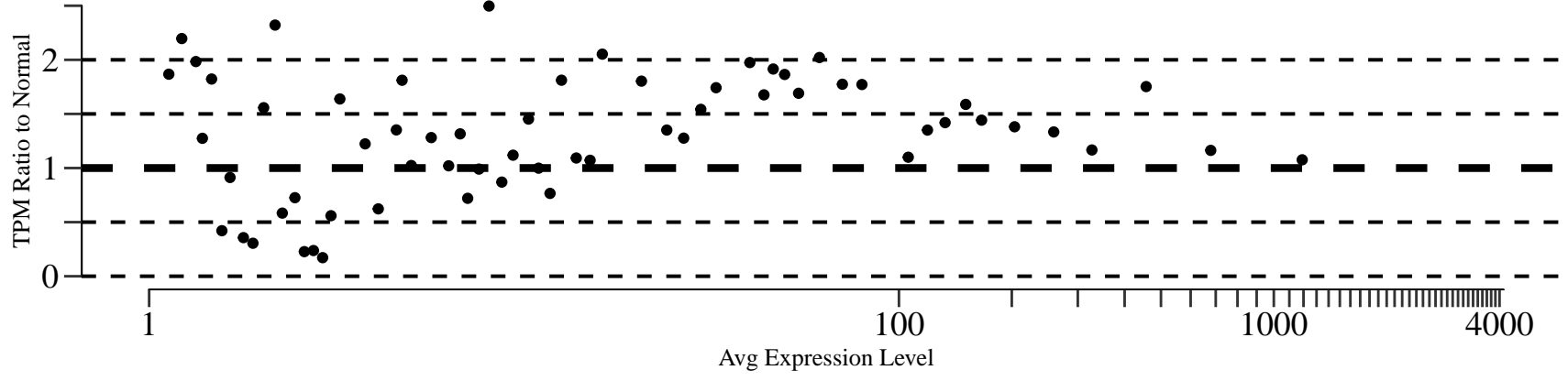
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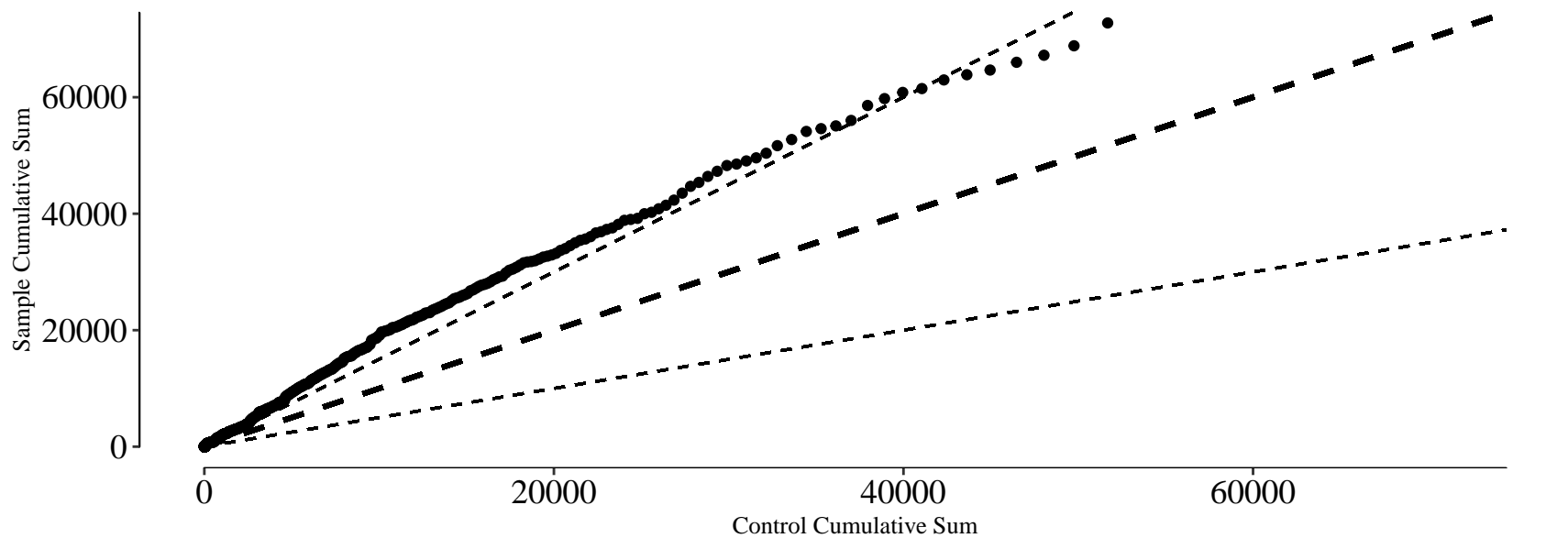
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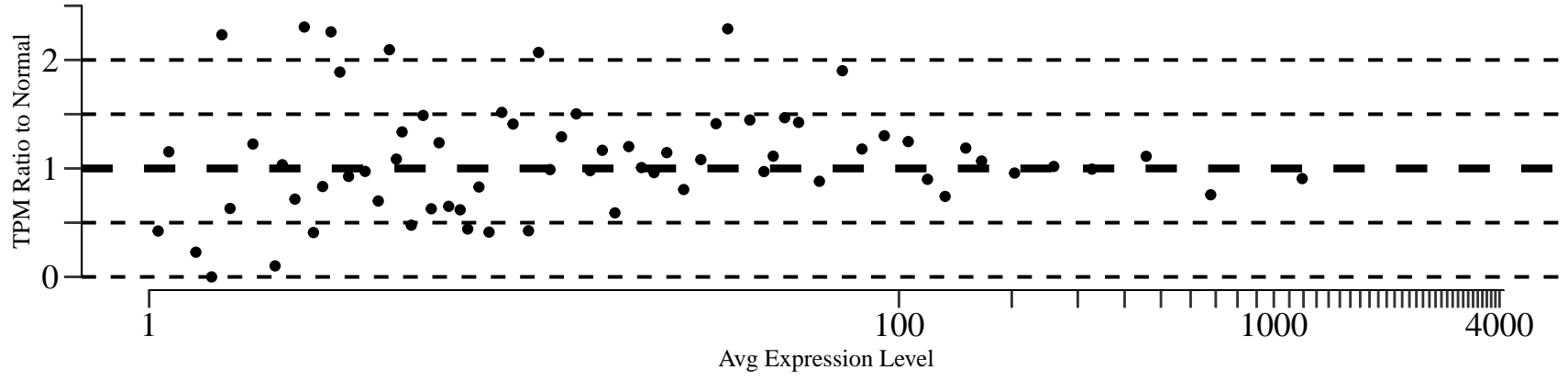
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ID: F227.2 | Chr: chr19



ID: F227.3 | Chr: chr19



ID: F227.3 | Chr: chr19

