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ASP: Chen and Huffman - Computational pipeline for whole-genome analysis of complex traits

[Introduction slides from May 11, 2017 All Hands call](#) [1]

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Award Type:	Working Group Analysis Support Program
Working Group or Committee:	
Analysis Center:	
Start Year:	2017

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Links

[1] https://topmed.nhlbi.nih.gov/system/files/meetings/Analysis_pipeline_presentation_20170511_ChenHuffman.pdf

[2] <https://topmed.nhlbi.nih.gov/project-group/dcc>