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## Mining for gold standard gene sets: maximizing the accuracy and utility of effector gene predictions from GWAS results

*Lists of the genes that are predicted to be effectors (causal genes) for complex traits and diseases are a valuable outcome of GWAS. How can we make them even more accessible and useful? Join our focus group to voice your opinion, learn about the results from our research into this area, and get a recap of our community workshop!*

