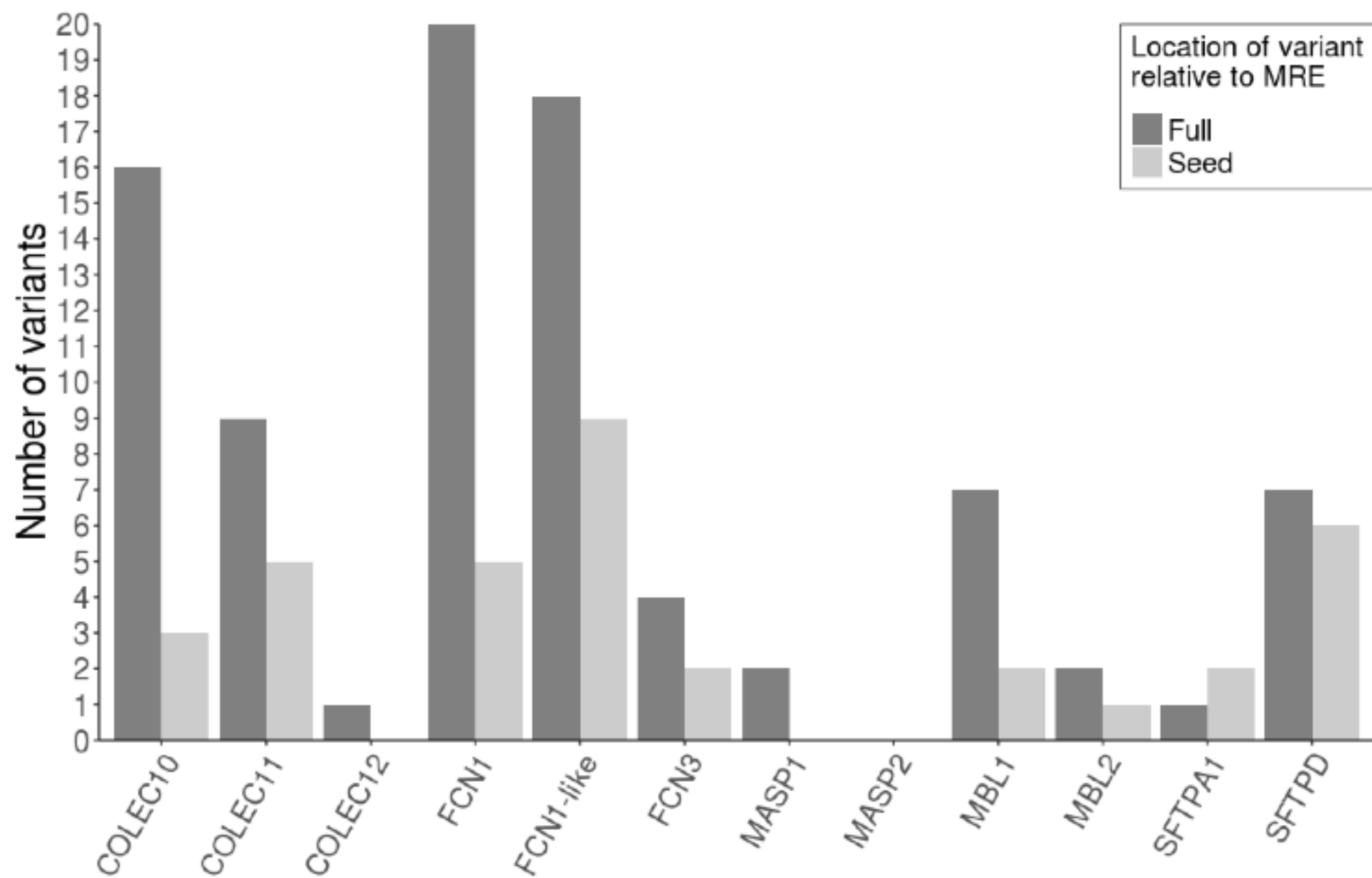






Predicted miRNA binding sites

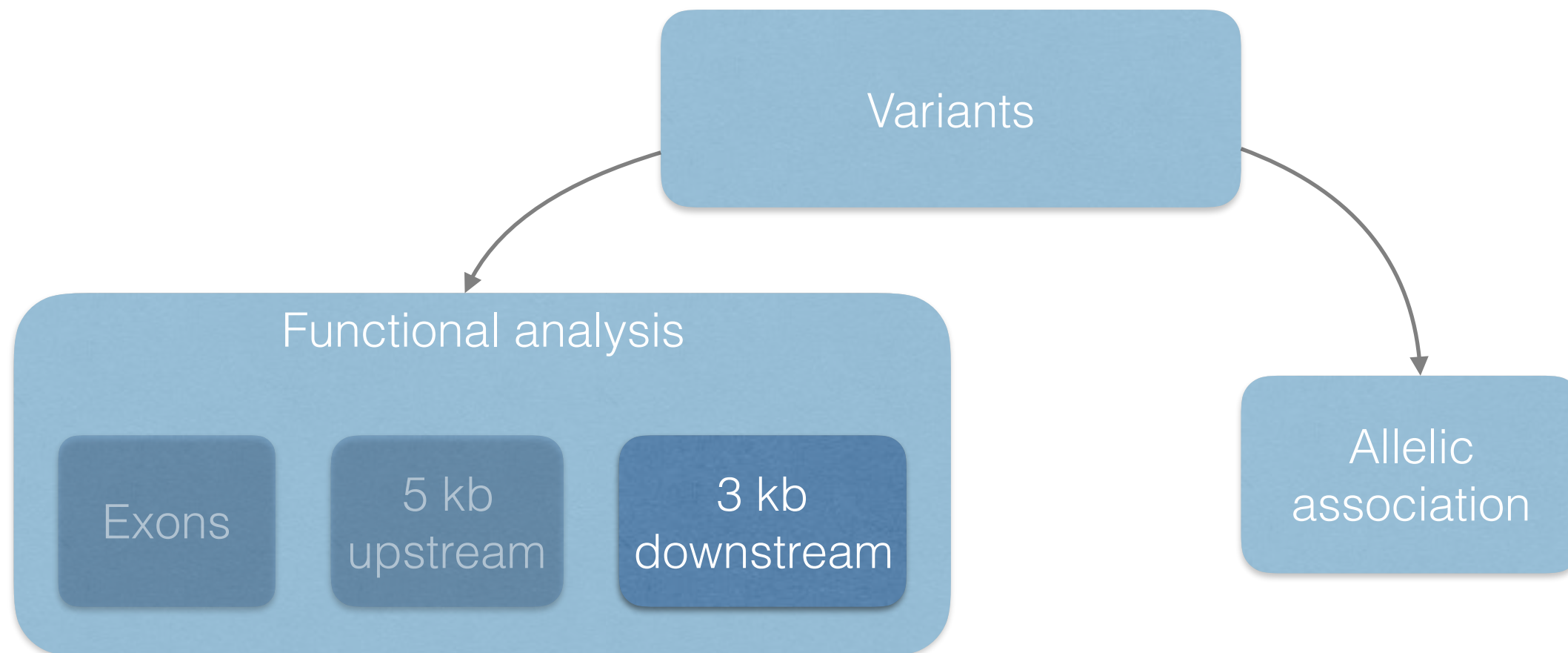


33 variants within the seed regions

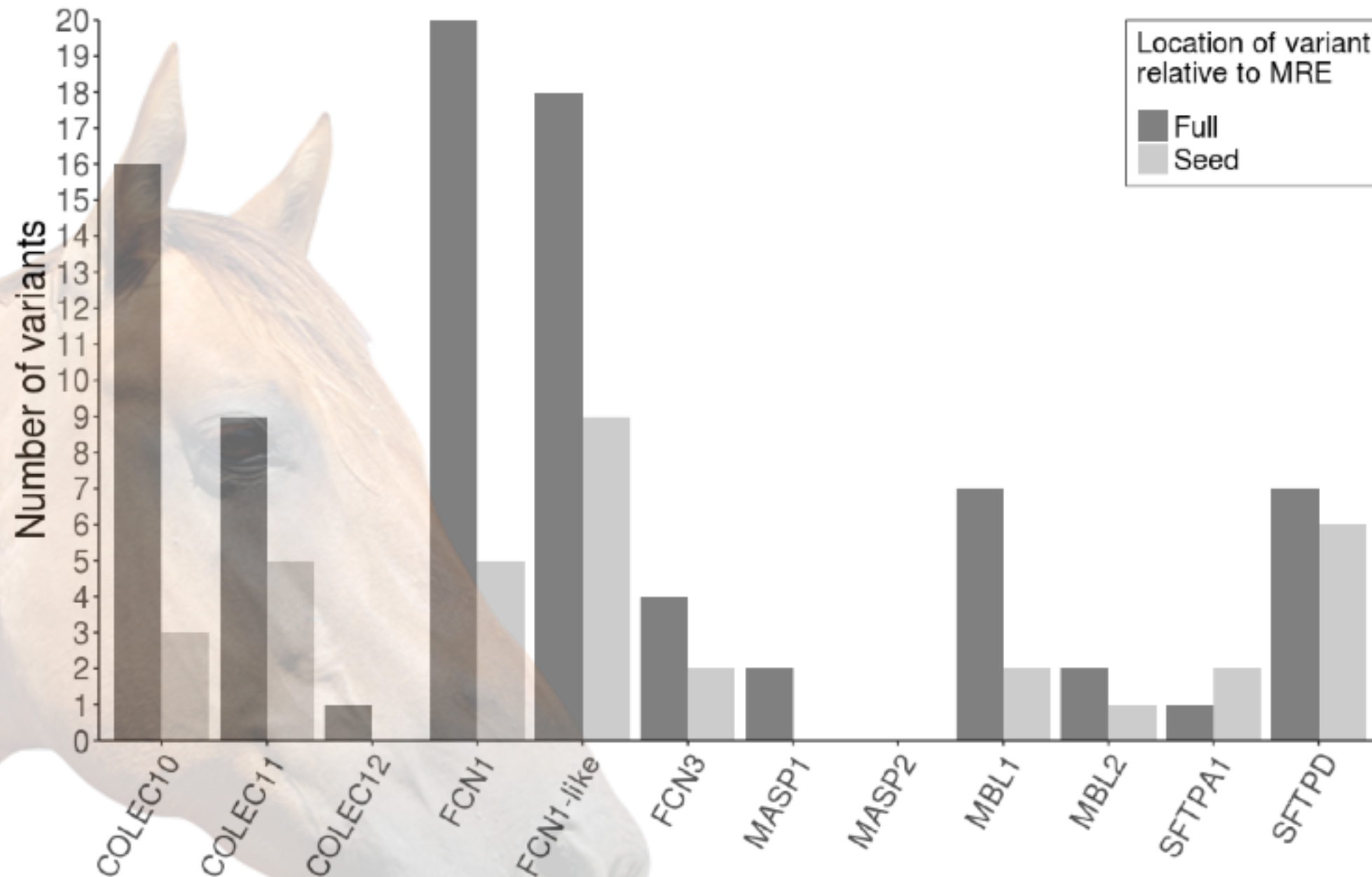


# Functional analysis and allelic association

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# Predicted miRNA binding sites



33 variants within the seed regions