

### SVAI\_combined\_sample

*With only localized variants*

**26**

*with 'hopping' to associated genes*

**547**

### Exome variants 1

*With only localized variants*

**26**

*with 'hopping' to associated genes*

**549**

### Exome Gene Dx

*With only localized variants*

**58**

*with 'hopping' to associated genes*

**885**