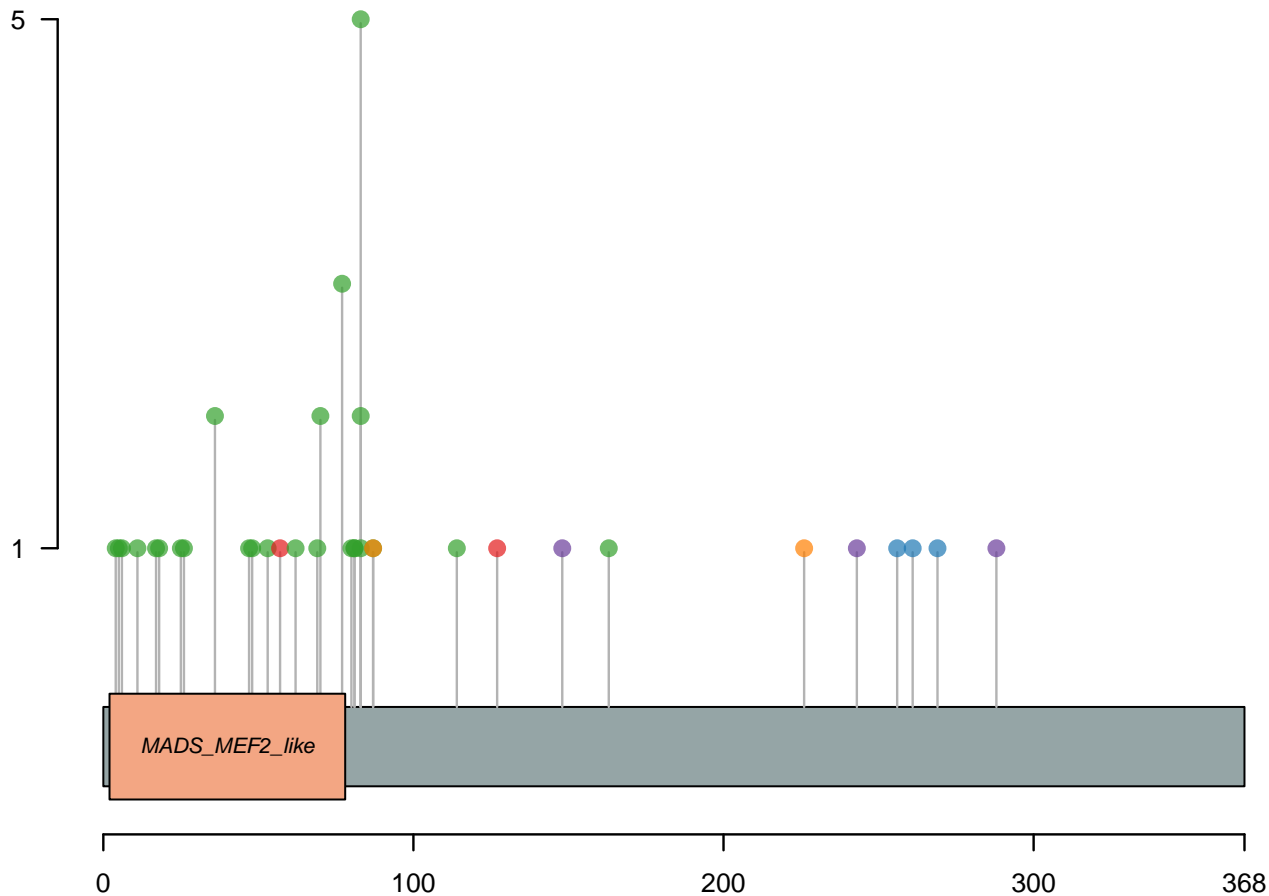# *MEF2B* : [Somatic Mutation Rate: 8.72%]
NM_001145785

Legend:
- Missense_Mutation
- Nonsense_Mutation
- Splice_Site
- Frame_Shift_Ins
- Frame_Shift_Del

Protein domain: MADS_MEF2_like

Axis: 0, 100, 200, 300, 368