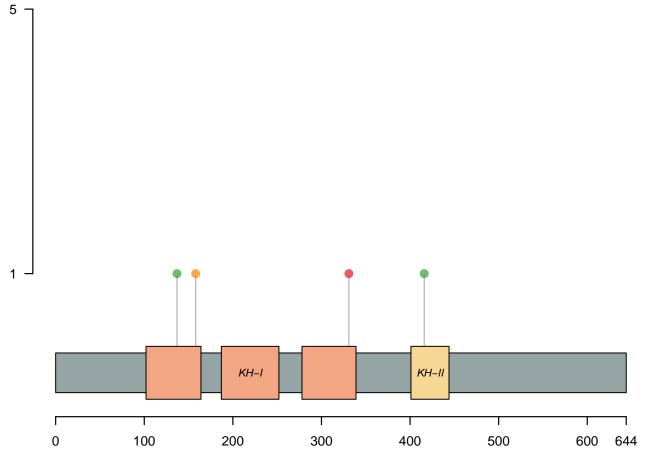
FUBP1 : [Somatic Mutation Rate: 1.06%] NM_003902



- Missense_Mutation
- Splice_Site
- Nonsense_Mutation