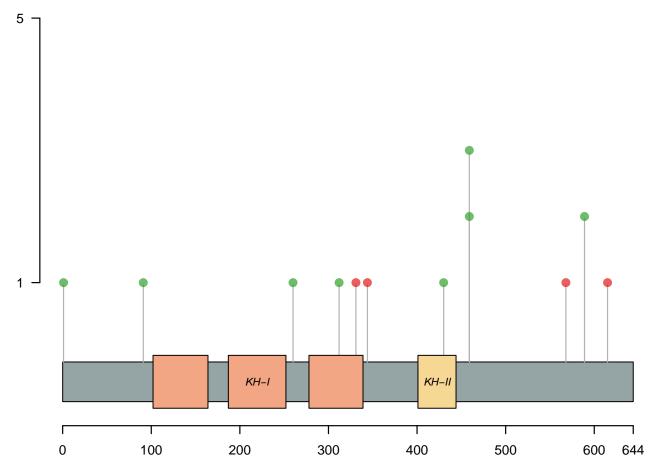
FUBP1 : [Somatic Mutation Rate: 1.85%] NM_003902



- Missense_Mutation
- Nonsense_Mutation