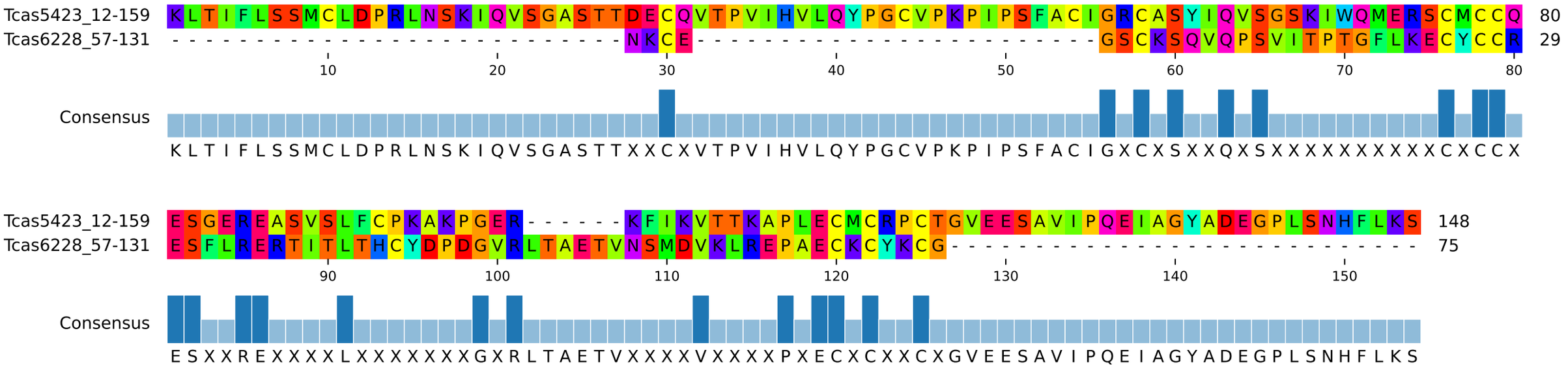
Tcas5423_12-159 KLTIFLSSMCLDPRLNSKIQVSGASTTDECQVTPVIHVLQYPGCVPKPIPSFACIGRCASYIQVSGSKIWQMERSCMCCQ 80
Tcas6228_57-131 ----------NKCE----------GSCKSQVQPSVITPTGFLKECYCCR 29
10 20 30 40 50 60 70 80
Consensus
KLTIFLSSMCLDPRLNSKIQVSGASTTXXCXVTPVIHVLQYPGCVPKPIPSFACIGXCXSXXQXSXXXXXXXXXXXCXCCX
Tcas5423_12-159 ESGERREASVSLFCPKAKPGER------KFIKVTTKAPLECMCRPCTGVEESAVIPQEIAGYADEGPLSNHFLKS 148
Tcas6228_57-131 ESFLRERTITLTHCYDPDGVRLTAETVNSMDVKLREPAECKCYKCG---------- 75
90 100 110 120 130 140 150
Consensus
ESXXREXXXXLXXXXXXXGXRLTAETVXXXXVXXXXPXECXCXXCXGVEESAVIPQEIAGYADEGPLSNHFLKS