

Scored residue / atom

CRAMP1 (300-800)

GON4L (1455-2024)

800 / 1455

2024

300 400 500 600 700 1555 1655 1755 1855 1955

CRAMP1 (300-800)

Aligned residue / atom

GON4L (1455-2024)

800 / 1455

1555 1655 1755 1855 1955 2024

