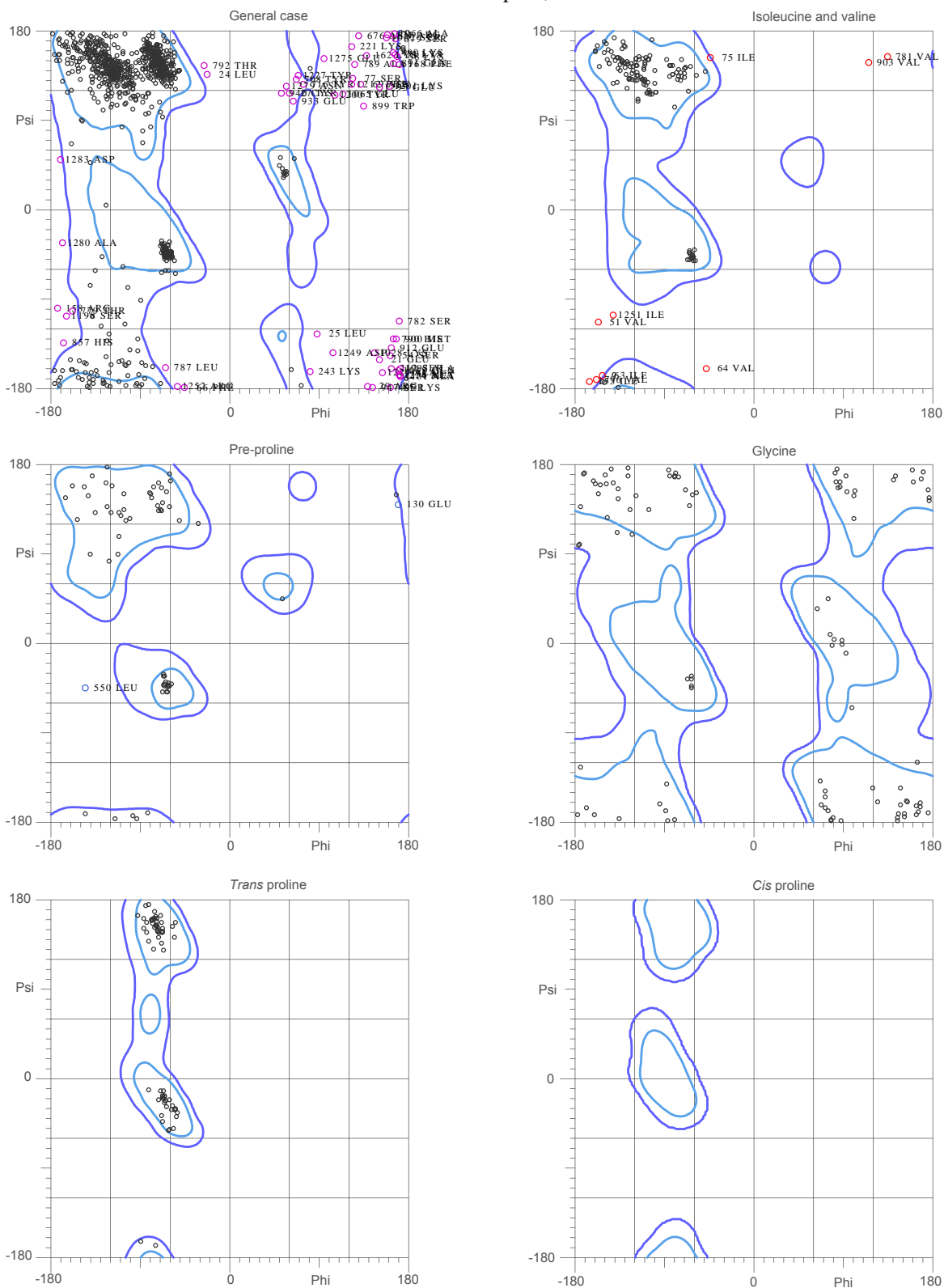


MolProbity Ramachandran analysis

6abu.B99990004.pdb, model 1



47.0% (10/21) of all residues were in favored (blue) regions.

96.0% (20/21) of all residues were in allowed (blue) regions.

None were in outlier (red) regions.

Residue (chain, residue number)

61 LEU (A, 117.0)

62 LEU (A, 118.0)

63 LEU (A, 119.0)

64 LEU (A, 120.0)

65 LEU (A, 121.0)

66 LEU (A, 122.0)

67 LEU (A, 123.0)

68 LEU (A, 124.0)

69 LEU (A, 125.0)

70 LEU (A, 126.0)

71 LEU (A, 127.0)

72 LEU (A, 128.0)

73 LEU (A, 129.0)

74 LEU (A, 130.0)

75 LEU (A, 131.0)

76 LEU (A, 132.0)

77 LEU (A, 133.0)

78 LEU (A, 134.0)

79 LEU (A, 135.0)

80 LEU (A, 136.0)

81 LEU (A, 137.0)

82 LEU (A, 138.0)

83 LEU (A, 139.0)

84 LEU (A, 140.0)

85 LEU (A, 141.0)

86 LEU (A, 142.0)

87 LEU (A, 143.0)

88 LEU (A, 144.0)

89 LEU (A, 145.0)

90 LEU (A, 146.0)

91 LEU (A, 147.0)

92 LEU (A, 148.0)

93 LEU (A, 149.0)

94 LEU (A, 150.0)

95 LEU (A, 151.0)

96 LEU (A, 152.0)

97 LEU (A, 153.0)

98 LEU (A, 154.0)

99 LEU (A, 155.0)

100 LEU (A, 156.0)

101 LEU (A, 157.0)

102 LEU (A, 158.0)

103 LEU (A, 159.0)

104 LEU (A, 160.0)