Using VQE with CVaR expectation value COBYLA optimzer 9 amino Bradykinin

April 4, 2023

```
[1]: from qiskit_research.protein_folding.interactions.random_interaction import (
         RandomInteraction,
     from qiskit_research.protein_folding.interactions.miyazawa_jernigan_interactionu
      →import (
         MiyazawaJerniganInteraction,
     from qiskit_research.protein_folding.peptide.peptide import Peptide
     from qiskit_research.protein_folding.protein_folding_problem import (
         ProteinFoldingProblem,
     )
     from qiskit_research.protein_folding.penalty_parameters import PenaltyParameters
     from qiskit.utils import algorithm_globals, QuantumInstance
     algorithm_globals.random_seed = 25
[2]: main_chain = "RPPGFSPFR" #Bradykinin Peptide
[3]: side_chains = [""] * 9
[4]: random_interaction = RandomInteraction()
     mj_interaction = MiyazawaJerniganInteraction()
[5]: penalty back = 10
     penalty_chiral = 10
     penalty_1 = 10
     penalty_terms = PenaltyParameters(penalty_chiral, penalty_back, penalty_1)
[6]: peptide = Peptide(main_chain, side_chains)
[7]: protein_folding_problem = ProteinFoldingProblem(peptide, mj_interaction,__
      →penalty_terms)
     qubit_op = protein_folding_problem.qubit_op()
```

[8]: print(qubit_op)

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5764.091 * IIIIIIIIIIIIII
+ 1240.0 * IIIIIIIIIIIIIIIIII
- 550.0 * IIIIIIIIIIIIIZZ
+ 550.0 * IIIIIIIIIIIIIZZZ
- 652.5 * IIIIIIIIIIIIIIIIII
- 652.5 * IIIIIIIIIIIIIIIIIIIII
- 652.5 * IIIIIIIIIIIIZZZZI
- 550.0 * IIIIIIIIIIIIIIII
- 550.0 * IIIIIIIIIIIIIIII
- 550.0 * IIIIIIIIIIZZZZIII
- 457.5 * IIIIIIIIIIIIII
- 457.5 * IIIIIIIIIIIIII
- 457.5 * IIIIIIIIZZZZIIIII
- 227.5 * IIIIIIZIZIIIIIII
- 227.5 * IIIIIIIIZIZIIIIIII
- 227.5 * IIIIIIZZZZIIIIIII
+ 690.0 * IIIIIIIIIIIIIIZZI
- 1250.0 * IIIIIIIIIIIIIIII
- 695.0 * IIIIIIIIIIIIZZIII
+ 460.0 * IIIIIIIIIIIIII
+ 555.0 * IIIIIIIIIIIIIII
- 555.0 * IIIIIIIIIIIIZZIIZ
- 904.5565 * IIIIIZIIIIIIIII
- 295.0 * IIIIIZIIIIIIIIZII
- 197.5 * IIIIIZIIIIIIIZZI
+ 302.5 * IIIIIZIIIIIIZIIII
+ 202.5 * IIIIIZIIIIIZZIII
+ 100.0 * IIIIIZIIIIIIZIZII
+ 100.0 * IIIIIZIIIIIIIZIZI
+ 100.0 * IIIIIZIIIIIIZZZZI
- 200.0 * IIIIIZIIIIIIIIIZ
+ 97.5 * IIIIIZIIIIIIIZZ
- 97.5 * IIIIIZIIIIIIIZZZ
- 100.0 * IIIIIZIIIIIIIZIIZ
+ 100.0 * IIIIIZIIIIIIZZIIZ
+ 230.0 * IIIIIIIIIIIIIIII
- 230.0 * IIIIIIIIIIIIIIIII
+ 2.5 * IIIIIZIIIIIIIIZI
- 2.5 * IIIIIZIIIIIIIZIII
+ 945.0 * IIIIIIIIIIIIIII
+ 490.0 * IIIIIIIIIIZZIIIII
+ 555.0 * IIIIIIIIIIIIIIIIII
+ 555.0 * IIIIIIIIIIIIIIIIIII
+ 555.0 * IIIIIIIIIIZZIIZZI
- 455.0 * IIIIIIIIIIIIIII
+ 455.0 * IIIIIIIIIIZZIIIIZ
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- + 7.5 * IIIIIZIIIIZIIIII
- + 5.0 * IIIIIZIIIIZZIIIII
- + 2.5 * IIIIIZIIIIZIII
- + 2.5 * IIIIIZIIIIIZI
- + 2.5 * IIIIIZIIIIZZIIZZI
- 2.5 * IIIIIZIIIIZIZIIII
- 2.5 * IIIIIZIIIIIZIZIII
- 2.5 * IIIIIZIIIIZZZZIII
- 2.5 * IIIIIZIIIIZ
- + 2.5 * IIIIIZIIIIZZIIIIZ
- 762.5 * IIIIIIIIIIIIII
- 400.0 * IIIIIIIIZZIIIIIII
- 462.5 * IIIIIIIIIIIIIIIII
- 462.5 * IIIIIIIIIIIIIIII
- 462.5 * IIIIIIIIZZIIIIZZI
- + 460.0 * IIIIIIIIIIIIIII
- + 460.0 * IIIIIIIIIIIIIII
- + 460.0 * IIIIIIIIZZIIZZIII
- + 362.5 * IIIIIIIIIIIIII
- 362.5 * IIIIIIIIZZIIIIIIZ
- 1616.925999999999 * IIIIZIIIIIIIII
- 400.0 * IIIIZIIIIIIIIIIII
- 267.5 * IIIIZIIIIIIIIZZI
- + 400.0 * IIIIZIIIIIIIZIIII
- + 267.5 * IIIIZIIIIIIIZZIII
- 400.0 * IIIIZIIIIIZIIIIII
- 267.5 * IIIIZIIIIIZZIIIII
- + 407.5 * IIIIZIIIZIIIIIII
- + 272.5 * IIIIZIIIZZIIIIIII
- + 132.5 * IIIIZIIIIIIIZIZII
- 132.5 * IIIIZIIIIIZIIIZII
- + 135.0 * IIIIZIIIZIIIIIZII
- + 132.5 * IIIIZIIIIIIIIZIZI
- 132.5 * IIIIZIIIIIIZI + 135.0 * IIIIZIIIIZI
- + 132.5 * IIIIZIIIIIIIZZZZI
- 132.5 * IIIIZIIIIIZZIIZZI
- + 135.0 * IIIIZIIIZZIIIIZZI
- + 132.5 * IIIIZIIIIIZIZIIII
- 135.0 * IIIIZIIIZIIIZIIII
- + 132.5 * IIIIZIIIIIIZIZIII
- 135.0 * IIIIZIIIIZIII
- + 132.5 * IIIIZIIIIIZZZZIII
- 135.0 * IIIIZIIIZZIIZZIII + 135.0 * IIIIZIIIZIZIIIIII
- + 135.0 * IIIIZIIIIZIZIIIII
- + 135.0 * IIIIZIIIZZZZIIIII
- 270.0 * IIIIZIIIIIIIIIZ

- + 132.5 * IIIIZIIIIIIIIZZ
- 132.5 * IIIIZIIIIIIIIZZZ
- 132.5 * IIIIZIIIIIIIIZIIZ
- + 132.5 * IIIIZIIIIIIIZZIIZ
- + 132.5 * IIIIZIIIIIZ
- 132.5 * IIIIZIIIIIZZIIIIZ
- 135.0 * IIIIZIIIIZIIIIIZ
- + 135.0 * IIIIZIIIZZIIIIIIZ
- + 235.0 * IIIIIIIIIIIIIII
- 130.0 * IIIIIIIIIIIIII
- + 2.5 * IIIIZIIIIIIIIIZI
- 2.5 * IIIIZIIIIIIIIIIII
- + 2.5 * IIIIZIIIIIIZIIIII
- 2.5 * IIIIZIIIIZIIIIII
- + 257.5 * IIIIIIZIIIIIIII
- + 130.0 * IIIIIIZZIIIIIIII
- + 232.5 * IIIIIIZIIIIIIIZII
- + 232.5 * IIIIIIIZIIIIIIIZI
- + 232.5 * IIIIIIZZIIIIIIZZI
- 230.0 * IIIIIIZIIIIIZIIII
- 230.0 * IIIIIIIZIIIIIZIII
- 230.0 * IIIIIIZZIIIIZZIII
- + 230.0 * IIIIIIZIIIZIIIIII
- + 230.0 * IIIIIIIZIIIZIIII
- + 230.0 * IIIIIIZZIIZZIIIII
- 127.5 * IIIIIIIIIIIII
- + 127.5 * IIIIIIZZIIIIIIIIZ
- + 7.5 * IIIIZIZIIIIIIIII
- + 5.0 * IIIIZIZZIIIIIIII
- + 2.5 * IIIIZIZIIIIIIIZII
- + 2.5 * IIIIZIIZIIIIIIIZI
- + 2.5 * IIIIZIZZIIIIIIZZI
- 2.5 * IIIIZIZIIIIIZIIII
- 2.5 * IIIIZIIZIIIIIZIII
- 2.5 * IIIIZIZZIIIIZZIII
- + 2.5 * IIIIZIZIIIZIIIII
- + 2.5 * IIIIZIIZIIIZIIII
- + 2.5 * IIIIZIZZIIZZIIIII - 2.5 * IIIIZIZIZIIIIIIII
- 2.5 * IIIIZIIZIZIIIIIII
- 2.5 * IIIIZIZZZZIIIIII
- 2.5 * IIIIZIIZIIIIIIIZ
- + 2.5 * IIIIZIZZIIIIIIIZ
- 686.949000000001 * IIIZIIIIIIIIII
- 190.0 * IIIZIIIIIIIIIIIII
- 100.0 * IIIZIIIIIIIIIIII
- 95.0 * IIIZIIIIIIIIIZZI
- + 190.0 * IIIZIIIIIIIIIIII

- + 100.0 * IIIZIIIIIIIIIIIII
- + 95.0 * IIIZIIIIIIIZZIII
- 195.0 * IIIZIIIIIIZIIIII
- 102.5 * IIIZIIIIIIIZIIIII
- 97.5 * IIIZIIIIIIZZIIIII
- + 95.0 * IIIZIIIIIIIIZIZII
- 97.5 * IIIZIIIIIIZIIIZII
- + 95.0 * IIIZIIIIIIIIIZIZI
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- + 95.0 * IIIZIIIIIIIZZZZI
- 97.5 * IIIZIIIIIIZZIIZZI
- + 97.5 * IIIZIIIIIIZIZIIII
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- + 97.5 * IIIZIIIIIIZZZZIII
- + 95.0 * IIIZIIIIIIIIIZZ
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- + 97.5 * IIIZIIIIIIIZIIIIZ
- 97.5 * IIIZIIIIIIZZIIIIZ
- + 5.0 * IIIZIIIIIIIIIZ
- 5.0 * IIIZIIIIZIIIIIII
- 2.5 * IIIZIIIIIZIIIIII
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- 2.5 * IIIZIIIIIZIIIIIZI
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- 2.5 * IIIZIIIIZZZZIIIII
- + 2.5 * IIIZIIIIIZIIIIIZ
- 2.5 * IIIZIIIIZZIIIIIIZ
- + 140.0 * IIIIIIIIIIIII
- 1344.53999999999 * IIZIIIIIIIIIII
- 265.0 * IIZIIIIIIIIIIIIII
- 137.5 * IIZIIIIIIIIIIIII
- 132.5 * IIZIIIIIIIIIIZZI
- + 265.0 * IIZIIIIIIIIIIIII
- + 137.5 * IIZIIIIIIIIIIIIII
- + 132.5 * IIZIIIIIIIIIZZIII
- 265.0 * IIZIIIIIIIIIIII
- 137.5 * IIZIIIIIIIIIIIII
- 132.5 * IIZIIIIIIIZZIIIII + 265.0 * IIZIIIIIZIIIIIIII
- + 137.5 * IIZIIIIIIZIIIIII

- + 132.5 * IIZIIIIIZZIIIIII
- 270.0 * IIZIIIZIIIIIIIII
- 140.0 * IIZIIIIZIIIIIIII
- 135.0 * IIZIIIZZIIIIIIII
- + 132.5 * IIZIIIIIIIIIIZIZII
- 132.5 * IIZIIIIIIIZIIIZII
- + 132.5 * IIZIIIIIZIIIIIZII
- 135.0 * IIZIIIZIIIIIIIZII
- + 132.5 * IIZIIIIIIIIIIIIIIII
- 132.5 * IIZIIIIIIIIZIIIZI
- + 132.5 * IIZIIIIIIZI
- 135.0 * IIZIIIIZIIIIIIIZI
- + 132.5 * IIZIIIIIIIIIZZZZI
- 132.5 * IIZIIIIIIIZZIIZZI
- + 132.5 * IIZIIIIIZZIIIIZZI
- 135.0 * IIZIIIZZIIIIIIZZI
- + 132.5 * IIZIIIIIIIIZIZIIII
- 132.5 * IIZIIIIIZIIIZIIII
- + 135.0 * IIZIIIZIIIIIZIIII
- + 132.5 * IIZIIIIIIIIIZIZIII
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- + 135.0 * IIZIIIIZIIIIZIII
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- + 135.0 * IIZIIIZZIIIIZZIII
- + 132.5 * IIZIIIIIZIZIIIIII
- 135.0 * IIZIIIZIIIZIIIII
- + 132.5 * IIZIIIIIIZIZIIIII
- 135.0 * IIZIIIIZIIIZIIIII
- + 132.5 * IIZIIIIIZZZZIIIII
- 135.0 * IIZIIIZZIIZZIIIII
- + 135.0 * IIZIIIZIZIIIIIII
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- 132.5 * IIZIIIIIIIZZIIIIZ
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- + 132.5 * IIZIIIIIZZIIIIIIZ
- + 135.0 * IIZIIIIZIIIIIIIIZ - 135.0 * IIZIIIZZIIIIIIII
- + 5.0 * IIZIIIIIIIIIIZ
- 591.696 * IZIIIIIIIIIIII
- + 95.0 * IZIIIIIIIIIIZIZII
- 95.0 * IZIIIIIIIIZIIIZII

- + 97.5 * IZIIIIIIZIIIIZII
- 95.0 * IZIIIIIIIIIIIIII
- + 95.0 * IZIIIIIIIIIIIZIZI
- 95.0 * IZIIIIIIIIIZIIIZI
- + 97.5 * IZIIIIIIIZI
- + 95.0 * IZIIIIIIIIIZZZZI
- 95.0 * IZIIIIIIIIZZIIZZI
- + 97.5 * IZIIIIIIZZIIIIZZI
- + 95.0 * IZIIIIIIIIZIZIIII
- 97.5 * IZIIIIIIZIIIZIIII
- + 95.0 * IZIIIIIIIIIIIIII
- + 95.0 * IZIIIIIIIIIZIZIII
- 97.5 * IZIIIIIIIZIIIZIII
- + 95.0 * IZIIIIIIIIZZZZIII
- 97.5 * IZIIIIIIZZIIZZIII
- + 97.5 * IZIIIIIIZIZIIIIII
- 95.0 * IZIIIIIIIIIIIII
- + 97.5 * IZIIIIIIIZIZIIIII
- + 97.5 * IZIIIIIIZZZZIIIII
- + 97.5 * IZIIIIIIZIIIIIII
- + 97.5 * IZIIIIIIIIIIZZ
- 97.5 * IZIIIIIIIIIIIZZZ
- 97.5 * IZIIIIIIIIIIZIIZ
- + 97.5 * IZIIIIIIIIIZZIIZ
- + 97.5 * IZIIIIIIIIIZIIIIZ
- 97.5 * IZIIIIIIIIZZIIIIZ
- 100.0 * IZIIIIIIIZIIIIIZ
- + 100.0 * IZIIIIIIZZIIIIIIZ
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- + 2.5 * IZIIIIIIIIIIIZZI
- 2.5 * IZIIIIIIIIIIIIIII
- 2.5 * IZIIIIIIIIIIZZIII
- + 2.5 * IZIIIIIIIIIIIII
- + 2.5 * IZIIIIIIIIZZIIIII
- 2.5 * IZIIIIIIIZIIIIII
- 2.5 * IZIIIIIIZZIIIIIII
- + 2.5 * IZIIIIZIIIIIIIII
- + 2.5 * IZIIIIZIIIIIIIZII
- + 2.5 * IZIIIIIZIIIIIIZI
- + 2.5 * IZIIIIZZIIIIIIZZI
- 2.5 * IZIIIIZIIIIIZIIII
- 2.5 * IZIIIIIZIIIIZIII
- 2.5 * IZIIIIZZIIIIZZIII
- + 2.5 * IZIIIIZIIIZIIIIII
- + 2.5 * IZIIIIIZIIIZIIII
- + 2.5 * IZIIIIZZIIZZIIIII
- 2.5 * IZIIIIZIZIIIIIII
- 2.5 * IZIIIIIZIZIIIIII

- 2.5 * IZIIIIZZZZIIIIII
- 2.5 * IZIIIIIZIIIIIIIZ
- + 2.5 * IZIIIIZZIIIIIIIZ
- 606.9235 * ZIIIIIIIIIIIII
- + 100.0 * ZIIIIIIIIIIIZIZII
- 100.0 * ZIIIIIIIIIZIIIZII
- + 100.0 * ZIIIIIIIZIIIIIZII
- 102.5 * ZIIIIIZIIIIIIIZII
- + 100.0 * ZIIIIIIIIIIIZIZI
- 100.0 * ZIIIIIIIIIIZIIIZI
- + 100.0 * ZIIIIIIIIZI
- 102.5 * ZIIIIIIZIIIIIIIZI
- + 100.0 * ZIIIIIIIIIIZZZZI
- 100.0 * ZIIIIIIIIIZZIIZZI
- + 100.0 * ZIIIIIIIZZIIIIZZI
- 102.5 * ZIIIIIZZIIIIIIZZI
- + 97.5 * ZIIIIIIIIIZIZIIII
- 97.5 * ZIIIIIIIZIIIZIIII
- + 100.0 * ZIIIIIZIIIIIZIIII
- + 97.5 * ZIIIIIIIIIIZIZIII
- 97.5 * ZIIIIIIIIZIIIZIII
- + 100.0 * ZIIIIIIZIIII
- + 97.5 * ZIIIIIIIIIZZZZIII
- 97.5 * ZIIIIIIIZZIIZZIII
- + 100.0 * ZIIIIIZZIIIIZZIII
- + 97.5 * ZIIIIIIIZIZIIIIII
- 100.0 * ZIIIIIZIIIZIIIII
- + 97.5 * ZIIIIIIIIZIZIIIII
- 100.0 * ZIIIIIIZIIIZIIIII
- + 97.5 * ZIIIIIIIZZZZIIIII
- 100.0 * ZIIIIIZZIIZZIIIII
- + 100.0 * ZIIIIIZIZIIIIIII
- + 100.0 * ZIIIIIIZIZIIIIIII
- + 100.0 * ZIIIIIZZZZIIIIII
- + 2.5 * ZIIIIIIIIIIIIIIIII
- 2.5 * ZIIIIIIIIIIIIIIII
- + 2.5 * ZIIIIIIIIIIIIIII
- 2.5 * ZIIIIIIIZIIIIIII
- + 2.5 * ZIIIIIZIIIIIIIII - 2.5 * ZIIIIIIIIIIIIZZ
- + 2.5 * ZIIIIIIIIIIIIIZZZ
- + 2.5 * ZIIIIIIIIIIIIZIIZ
- 2.5 * ZIIIIIIIIIIZZIIZ
- 2.5 * ZIIIIIIIIIIIIII
- + 2.5 * ZIIIIIIIIIZZIIIIZ
- + 2.5 * ZIIIIIIIIZIIIIIZ
- 2.5 * ZIIIIIIIZZIIIIIIZ
- 2.5 * ZIIIIIIZIIIIIIIZ

+ 2.5 * ZIIIIIZZIIIIIIIZ

```
[9]: # Get the qubit operator and count the number of qubits
      qubit_op = protein_folding_problem.qubit_op()
      num_qubits = qubit_op.num_qubits
      print(f"The protein folding problem requires {num qubits} qubits.")
     The protein folding problem requires 17 qubits.
[10]: from qiskit.circuit.library import RealAmplitudes
      from qiskit.algorithms.optimizers import COBYLA, SPSA
      from qiskit.algorithms import NumPyMinimumEigensolver
      from qiskit.algorithms.minimum_eigensolvers import SamplingVQE
      from qiskit import execute, Aer
      from qiskit.primitives import Sampler
[11]: # set classical optimizer
      optimizer = SPSA(maxiter=150)
      # set variational ansatz
      ansatz = RealAmplitudes(reps=1)
      counts = \Pi
      values = []
[12]: def store_intermediate_result(eval_count, parameters, mean, std):
          counts.append(eval_count)
          values.append(mean)
      # initialize VQE using CVaR with alpha = 0.1
      vqe = SamplingVQE(
          Sampler(),
          ansatz=ansatz,
          optimizer=optimizer,
          aggregation=0.1,
          callback=store intermediate result,
      raw_result = vqe.compute_minimum_eigenvalue(qubit_op)
      print(raw_result)
     SamplingMinimumEigensolverResult:
             Eigenvalue: 59.883826428389135
             Best measurement
     : {'state': 69827, 'bitstring': '10001000011000011', 'value':
     (-2.301000000000446+0j), 'probability': 1.345470155e-05}
```

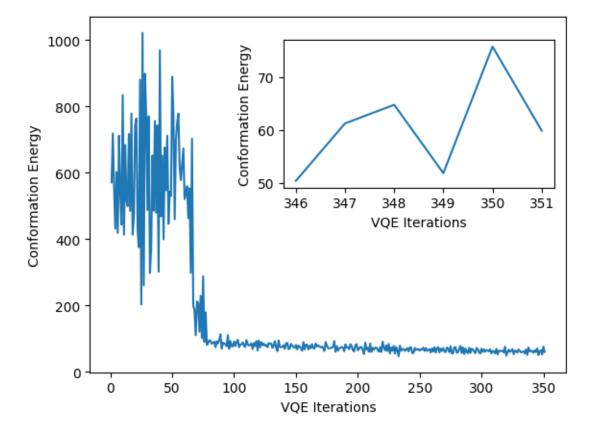
```
[19]: import matplotlib.pyplot as plt

fig = plt.figure()

plt.plot(counts, values)
plt.ylabel("Conformation Energy")
plt.xlabel("VQE Iterations")

fig.add_axes([0.44, 0.51, 0.44, 0.32])

plt.plot(counts[345:], values[345:])
plt.ylabel("Conformation Energy")
plt.xlabel("VQE Iterations")
plt.show()
```

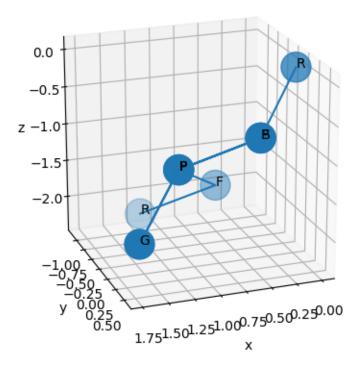


```
[14]: result = protein_folding_problem.interpret(raw_result=raw_result)
print(
    "The bitstring representing the shape of the protein during optimization is:
    ",
    result.turn_sequence,
)
```

```
print("The expanded expression is:", result.get_result_binary_vector())
     The bitstring representing the shape of the protein during optimization is:
     01010000010000100
     The expanded expression is: 0_____1___0_1___0_0_
     _____0001000010_0___
[15]: print(
         f"The folded protein's main sequence of turns is: {result.
      →protein_shape_decoder.main_turns}"
     print(f"and the side turn sequences are: {result.protein_shape_decoder.

side_turns}")
     The folded protein's main sequence of turns is: [1, 0, 1, 1, 0, 0, 2, 0]
     and the side turn sequences are: [None, None, None, None, None, None, None,
     None, None]
[16]: print(result.protein_shape_file_gen.get_xyz_data())
     [['R' '0.0' '0.0' '0.0']
      ['P' '0.5773502691896258' '0.5773502691896258' '-0.5773502691896258']
      ['P' '1.1547005383792517' '0.0' '-1.1547005383792517']
      ['G' '1.7320508075688776' '0.5773502691896258' '-1.7320508075688776']
      ['F' '1.154700538379252' '0.0' '-1.154700538379252']
      ['S' '0.5773502691896261' '0.5773502691896258' '-0.5773502691896261']
      ['P' '1.154700538379252' '0.0' '-1.154700538379252']
      ['F' '0.5773502691896261' '-0.5773502691896258' '-1.7320508075688776']
      ['R' '1.154700538379252' '-1.1547005383792517' '-2.3094010767585034']]
[20]: fig = result.get_figure(title="Protein Structure of Bradykinin Peptide", u
      →ticks=True, grid=True)
     fig.get_axes()[0].view_init(20, 70)
```

Protein Structure of Bradykinin Peptide



[]: