

Fourier Spectral Analysis of Coding Sequences

Automated Report

November 20, 2025

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- 1 Group 1: Single Peak at 0.33
- 2 Group 2: Single Peak Elsewhere
- 3 Group 3: Multiple Peaks

3.1 SCFR_NC_060941.1_6087888_6094167

3.1.1 Protein: frame-1

Gene ID: SCFR

Nucleotide Accession: NC_060941.1

Sequence Length: 6279 bp

Protein Description:

Stop-codon-free region (6087888-6094167)

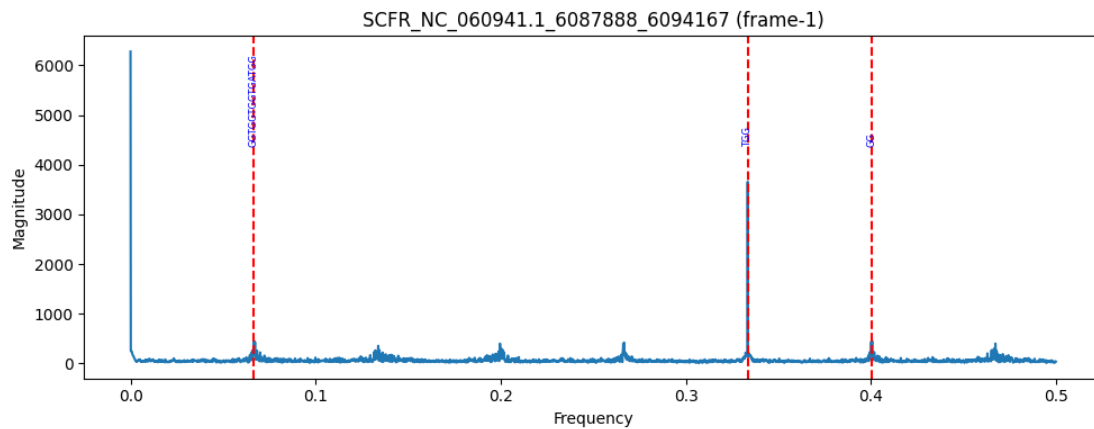


Figure 1: FFT Spectrum for frame-1

| Frequency | Motifs |
|-----------|----------------------------------------------------------------------------------------------------------|
| 0.333 | TGG (1154), GGT (1145), GTG (1145), GGG (681), ATG (484) |
| 0.401 | GG (1983), TG (1643), GT (1162), GA (636), AT (491) |
| 0.067 | GGTGGTGGTGGTGG (285), TGGTGATGGGGGTG (214), GGTGATGGGGGTGG (161), GTGATGGGGTGG (161), GTGATGGGGTGG (161) |

3.2 SCFR_NC_060941.1_81645697_81650701

3.2.1 Protein: frame-3

Gene ID: SCFR

Nucleotide Accession: NC_060941.1

Sequence Length: 5004 bp

Protein Description:

Stop-codon-free region (81645697-81650701)

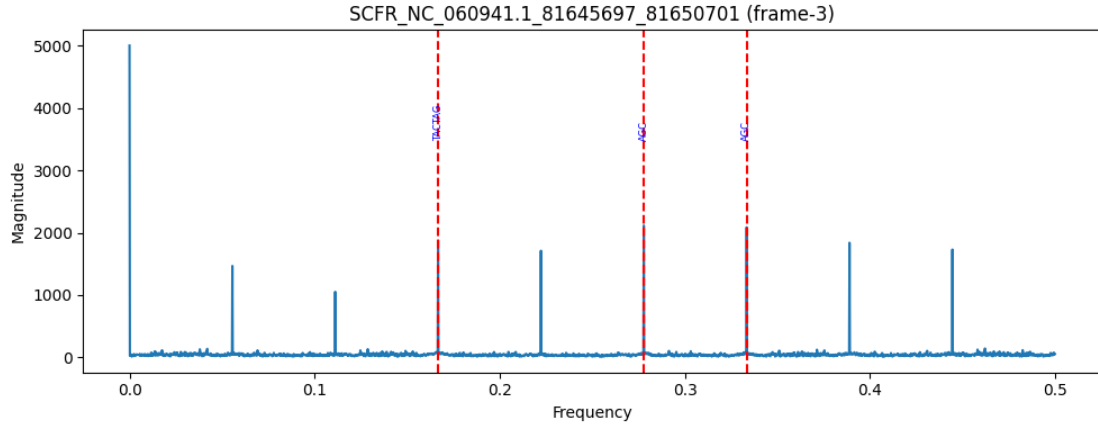


Figure 2: FFT Spectrum for frame-3

| Frequency | Motifs |
|-----------|----------------------------------------------------------------------|
| 0.278 | AGC (418), CTA (417), CTG (398), ACT (380), CCT (275) |
| 0.333 | AGC (418), CTA (417), CTG (398), ACT (380), CCT (275) |
| 0.167 | TACTAG (249), CTAGCA (249), ACTAGC (248), TAGCAC (241), CCTGGC (165) |

3.3 SCFR_NC_060941.1_6087991_6094060

3.3.1 Protein: frame-3

Gene ID: SCFR
Nucleotide Accession: NC_060941.1
Sequence Length: 6069 bp
Protein Description:
Stop-codon-free region (6087991-6094060)

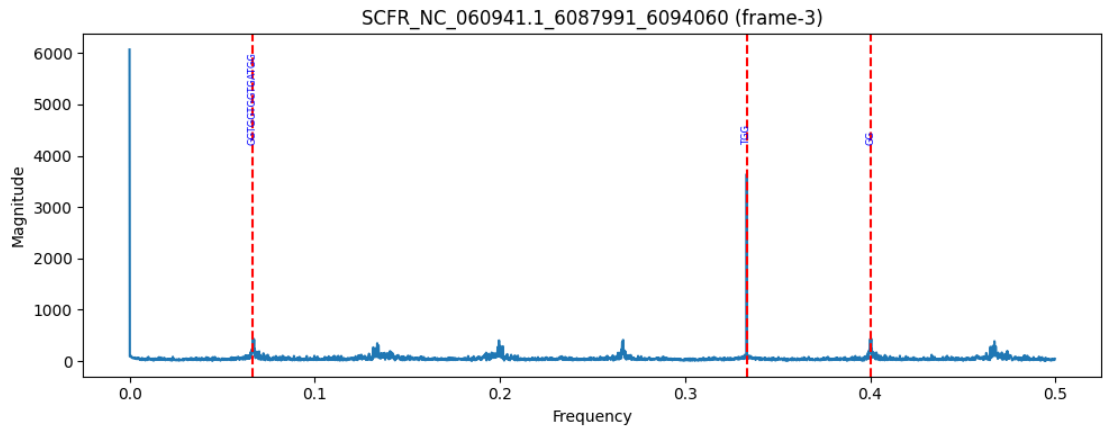


Figure 3: FFT Spectrum for frame-3

| Frequency | Motifs |
|-----------|------------------------------------------------------------------------------------------------------------|
| 0.333 | TGG (1149), GGT (1142), GTG (1142), GGG (673), ATG (480) |
| 0.067 | GGTGGTGGTGGTGG (285), TGGTGATGGGGGTG (214), GGTGATGGGGGTGG (161), GTGATGGGGTGG (128), GGTGGTGGTGGTGG (285) |
| 0.401 | GG (1963), TG (1628), GT (1154), GA (622), AT (482) |

3.4 SCFR_NC_060941.1_6087832_6094120

3.4.1 Protein: frame2

Gene ID: SCFR

Nucleotide Accession: NC_060941.1

Sequence Length: 6288 bp

Protein Description:

Stop-codon-free region (6087832-6094120)

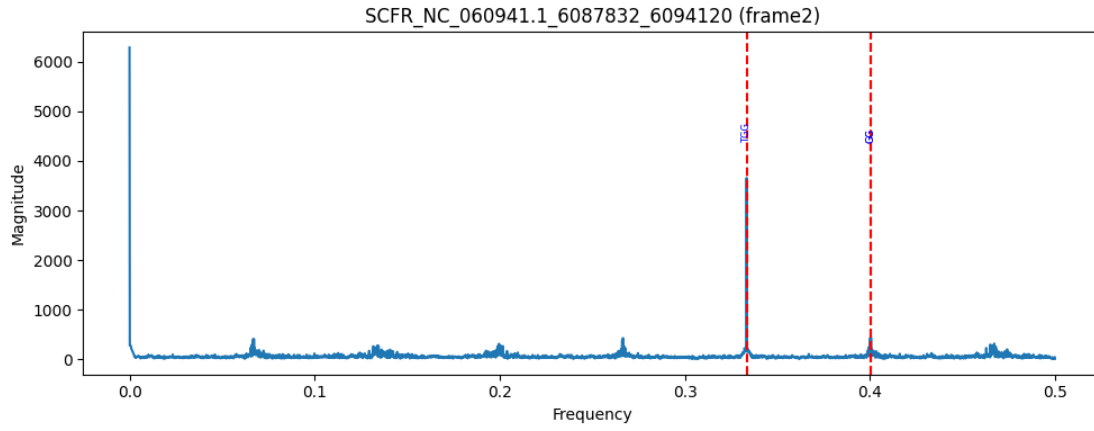


Figure 4: FFT Spectrum for frame2

| Frequency | Motifs |
|-----------|----------------------------------------------------------|
| 0.333 | TGG (1152), GGT (1145), GTG (1144), GGG (679), ATG (485) |
| 0.4 | GG (1980), TG (1640), GT (1161), GA (637), AT (492) |
| 0.401 | GG (1980), TG (1640), GT (1161), GA (637), AT (492) |

3.5 SCFR_NC_060941.1_6088040_6094049

3.5.1 Protein: frame-2

Gene ID: SCFR

Nucleotide Accession: NC_060941.1

Sequence Length: 6009 bp

Protein Description:

Stop-codon-free region (6088040-6094049)

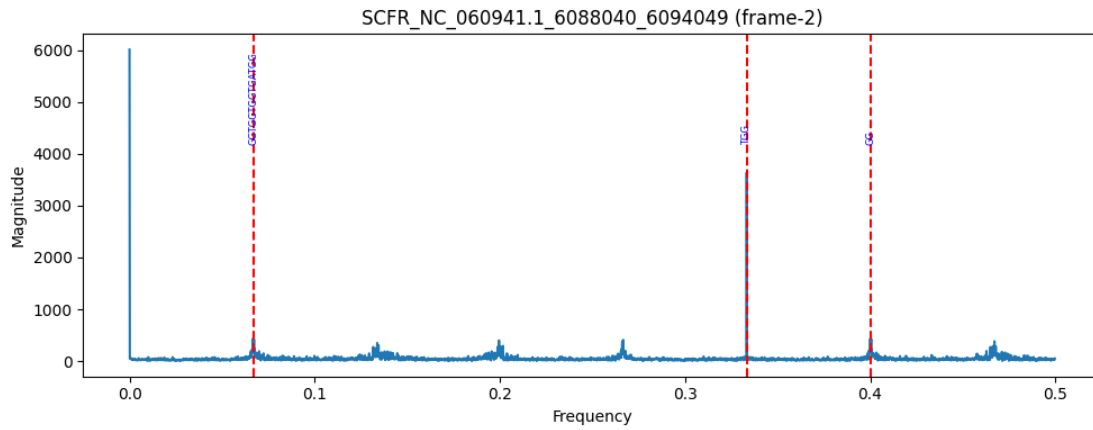


Figure 5: FFT Spectrum for frame-2

| Frequency | Motifs |
|-----------|----------------------------------------------------------------------------|
| 0.333 | TGG (1146), GGT (1142), GTG (1142), GGG (669), GAT (477) |
| 0.067 | GGTGGTGGTGATGG (285), TGGTGATGGGGGTG (214), GGTGATGGGGGTGG (161), GTGATGGG |
| 0.4 | GG (1954), TG (1624), GT (1154), GA (619), AT (477) |

3.6 SCFR_NC_060941.1_6088041_6094404

3.6.1 Protein: frame1

Gene ID: SCFR

Nucleotide Accession: NC_060941.1

Sequence Length: 6363 bp

Protein Description:

Stop-codon-free region (6088041-6094404)

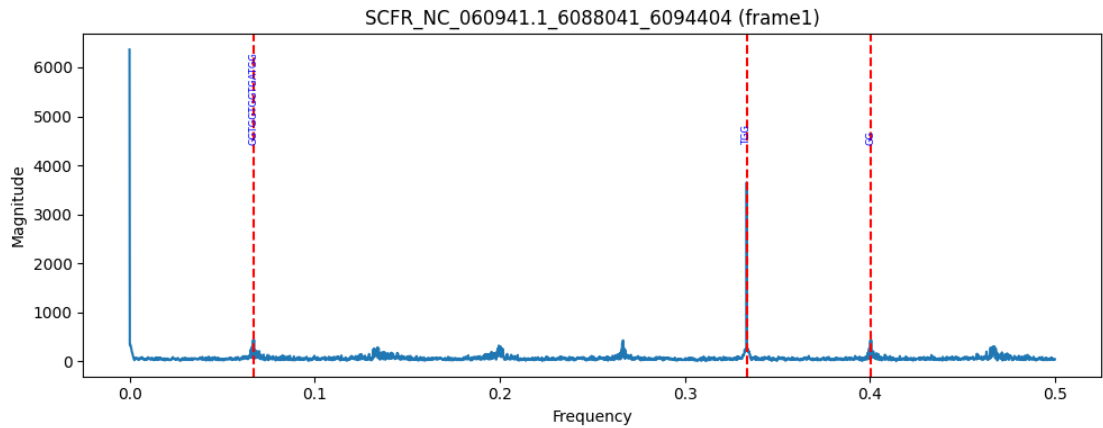


Figure 6: FFT Spectrum for frame1

| Frequency | Motifs |
|-----------|----------------------------------------------------------------------------|
| 0.333 | TGG (1157), GGT (1146), GTG (1146), GGG (673), ATG (483) |
| 0.4 | GG (1972), TG (1654), GT (1170), GA (630), AT (498) |
| 0.067 | GGTGGTGGTGATGG (285), TGGTGATGGGGGTG (214), GGTGATGGGGGTGG (161), GTGATGGG |