

Table 2 Standard reduction potentials (E°) of redox systems involved in biological electron transfer at pH 7. Adapted from Roehm *et al.*⁶²

| Redox species | E° (V) | n | Redox species | E° (V) | n |
|---|---------------|-----|---|---------------|-----|
| Ferredoxins | -0.27--0.5 | — | UQ/UQH ₂ | +0.06 | 2 |
| H ⁺ /H ₂ | -0.42 | 2 | UQ [•] /UQH ₂ | +0.19 | 1 |
| NADP ⁺ /NADPH | -0.32 | 2 | Cytochrome c ₁ (Fe ³⁺ /Fe ²⁺) | +0.22 | 1 |
| Lipoamide _{ox} /lipoamide _{red} | -0.29 | 2 | Cytochrome c (Fe ³⁺ /Fe ²⁺) | +0.25 | 1 |
| FMN/FMNH ₂ ^a | -0.20 | 2 | Riesk [2Fe-2S] (Fe ³⁺ /Fe ²⁺) | +0.28 | 1 |
| FAD/FADH ₂ ^a | -0.20 | 2 | Cytochrome a (Fe ³⁺ /Fe ²⁺) | +0.29 | 1 |
| Cytochrome b _L (Fe ³⁺ /Fe ²⁺) | -0.10 | 1 | Cytochrome a ₃ (Fe ³⁺ /Fe ²⁺) | +0.35 | 1 |
| FAD/FADH ₂ ^b | 0.0--0.1 | 2 | Cytochrome f (Fe ³⁺ /Fe ²⁺) | +0.37 | 1 |
| UQ/UQH [•] | +0.03 | 1 | O ₂ /H ₂ O | +0.82 | 2 |
| Cytochrome b _H (Fe ³⁺ /Fe ²⁺) | +0.06 | 1 | | | |

^a Free molecule. ^b Protein-bound.

62 K. H. Roehm, *eLS*, 2001, DOI: 10.1038/npg.els.0001373.