

Perspective

Cite

Morelli AM (2026) Putative Membrane Energy Reserves predicted by the Proton Capacitor Theory. BEC preprints 2026.1.

<https://doi.org/10.26124/bec.2026-0001>

Conflicts of interest

The author declares no conflict of interest.

Received 2026-01-29

Accepted 2026-01-29

Online 2026-02-10

Keywords

mitochondria;
chemiosmotic theory;
mitochondria derived vesicles;
evolution;
endosymbiotic theory

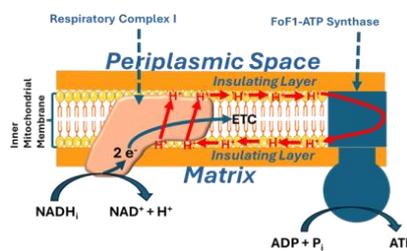
Putative Membrane Energy Reserves predicted by the Proton Capacitor Theory

 Alessandro Maria Morelli

University of Genoa (Italy) - Retired

Correspondence: morelliales@gmail.com

Summary



A significant number of reports indicate that oxidative phosphorylation (OXPHOS) occurs at multiple extra-mitochondrial sites, and this is not predicted by

canonical mitochondrial criteria. This requires an update of the endosymbiotic theory by considering that ancestral bacteria, in becoming mitochondria, retained the ability to vesiculate and widespread distribution in organism districts of mitochondria derived vesicles (MDVs), which possess active OXPHOS, confirms that mitochondrial vesiculation is an active process in eukaryotic cells. MDVs can fuse with other membrane systems, and the succession of these processes appears to be a route for the export of OXPHOS to many subcellular and extra-cellular districts. Considering that the chemiosmotic theory is not applicable to extra-mitochondrial districts, a new theory is needed. Here is presented the Proton Capacitor Theory that is independent of the parameters Membrane Potential and Protomotive Force which has led to exhaustive discussions in the past and therefore the modalities with which the aerobic synthesis of ATP occurs both in the mitochondria and outside the mitochondria are greatly simplified by the Proton Capacitor Theory also with regards to computational thermodynamics. The proton movement scheme proposed here allows us to hypothesize that i) membranes can have a buffering action towards protons and therefore could be sites of rapidly usable energy deposits here called Proton Capacitors, and ii) it is possible that OXPHOS can occur in two phases, i.e. "charging" and "discharging" of the proton capacitor, consistently with the hypothesis that underlies the recently formulated Sleep Theory.

1. Introduction

After an oxygen-poor atmosphere the amount of oxygen raised enormously due to the oxygen production by photosynthetic bacteria during the Great Oxidation Event around 2.4–2.1 billion years ago [1]. So, to develop reactions that consumed oxygen, according to the endosymbiotic theory [2], the aerobic bacteria were phagocytized by ancestral eukaryotes, but this statement needs to be updated because it is unlikely that bacteria, precursors of mitochondria, were engulfed by ancestral eukaryotes to supply the cell with ATP, a precious resource that would have helped these ancestral eukaryotes to even better phagocytose these "intruders". Some have rightly called it the "*Suicide Hypothesis*" [3].

It has been hypothesized [4] that bacteria, which possess oxidative phosphorylation (OXPHOS) but have poorly developed internal membranes (bacteria lack the cristae typical of mitochondria), in evolving into mitochondria received from ancestral eukaryotes [5] [6] the instructions for developing the surface of the internal membrane, i.e. the cristae, which in eukaryotes is highly developed in the form of endoplasmic reticulum. Thus a new hypothesis [4] prospects a true symbiosis, since bacteria transferred the OXPHOS machinery to the endoplasmic reticulum of ancestral eukaryotes, and ancestral eukaryotes transferred the instructions for highly developed internal membranes, i.e. the mitochondrial cristae, to bacteria which have become mitochondria.

This process which leads to the fusion of the mitochondrial membrane with other cellular membranes, endoplasmic reticulum first and foremost, means that the traditional OXPHOS mitochondrial functions -with coupled aerobic ATP synthesis- was expressed with high metabolite fluxes even in other cellular membranes and in fact it has been seen that OXPHOS is operative in vertebrate photoreceptors [7], myelin [8], platelets [9] plasma cells [10], sarcolemma of skeletal muscle [11] and everything suggests that extra-mitochondrial OXPHOS is operative in all subcellular districts of almost all living organisms.

This extra-mitochondrial diffusion of the OXPHOS which is considered to be exclusively mitochondrial requires the development of a new theory (here called "Proton Capacitor Theory") on the coupling between proton movement and ATP synthesis. The "proton capacitor" is a system for the storage of ready-to-use energy which has already been applied to a hypothesis on the mechanisms of sleep [12].

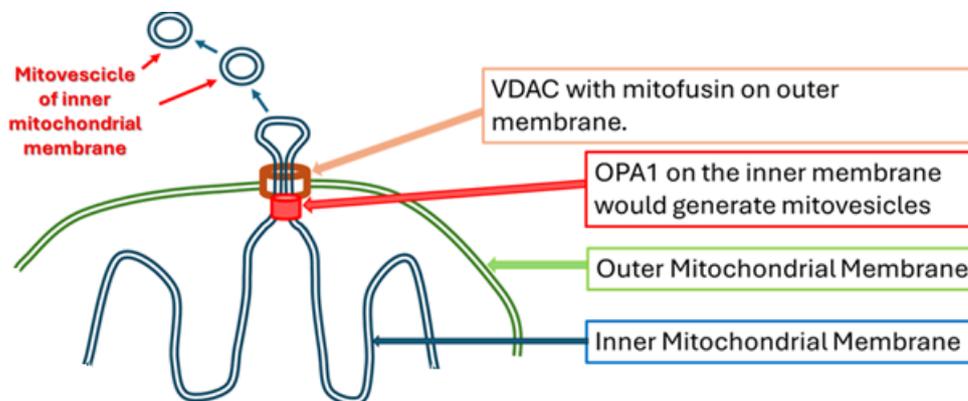
In developing this new theory, we will take into account the principle that chemical processes are much more efficient if they occur in a homogeneous phase, and therefore the movement of protons is located exclusively within the proteolipid phase of the membrane.

2. Production of Mitochondrial Derived Vesicles

An property of bacteria must be taken into account, namely their extreme versatility in producing vesicles [13], property that has been retained by mitochondria that possess a plurality of molecular devices for vesicle production, namely Mitofusin 1 and 2, OPA1, Drp1 [14]. Interestingly, their functional deficit causes serious neurological pathologies [15]. The production of mitochondria-derived vesicles (MDVs) in eukaryotes is a highly active and multifaceted process that has been the subject of

52 much research in recent years [16][17][18][19][20]. MDVs can fuse in the
 53 endoplasmic reticulum [21] where they form rafts that contain about 24% mitochondrial
 54 proteins [22]. The myelin sheath also contains rafts [23] [24] and this is consistent with
 55 the probable transfer of rafts to the oligodendrocyte endoplasmic reticulum (ER), which
 56 would then fuse with the forming myelin sheath.

57
 58 The study of MDVs thus opens up fascinating scenarios, particularly the visionary
 59 idea that mitovesicles bud from mitochondria to spread and fuse with endocellular
 60 membrane, endoplasmic reticulum *in primis*, where ATP synthesis can occur *in situ*
 61 thanks to this import of the OXPHOS molecular machinery. Fig. 1 shows a possible
 62 vesiculation process of the mitochondrial inner membrane that passes through the
 63 maxi-pore Voltage Dependent Anion Channel (VDAC) [25] present on the
 64 mitochondrial outer membrane.
 65



66
 67 **Figure 1. Proposal extrusion and delivery of mitovesicles of inner mitochondrial**
 68 **membrane fragments with "narrowing" by OPA1 inserted in inner mitochondrial**
 69 **membrane and extrusion through VDAC inserted in the outer mitochondrial membrane.**

70
 71 The VDAC, which is located in the outer membrane of the mitochondrion, plays
 72 an important role, as it may extrude the inner mitochondrial membrane to produce
 73 MDV. This hypothesis stems from the fact that VDAC is a protein with a structure of 19
 74 antiparallel beta-barrel strands arranged in a barrel shape with an internal diameter of
 75 3.15 nm [26] but it could tetramerize [27] forming new hydrogen bonds between one
 76 end of a VDAC with the end of another VDAC forming a ring with a diameter of ~13 nm,
 77 suitable for "squeezing" and extruding MDV.
 78

79 This vesiculation process would export the complex OXPHOS machinery to
 80 other sites, giving these sites the ability to synthesize ATP *in situ*.

81 All this goes beyond the traditional model that attributes the aerobic synthesis of
 82 ATP to mitochondria alone. Note that MDVs contain FoF1-ATP Synthase and
 83 synthesize ATP [28][29][30].
 84
 85
 86

3. Formulation of Proton Capacitor Theory

Since the chemiosmotic theory [31] is not applicable to non-mitochondrial membranes expressing OXPHOS, it is necessary to formulate a new theory which I currently call Proton Capacitor Theory. In doing so, previous reports that have addressed this topic are taken into account [32][33][34][35][36][19][37][38]. A recent article by S. Ji clearly identifies the limitations of the chemiosmotic theory [39]. Fig. 2A and 2B show, respectively, the scheme predicted by the chemiosmotic theory and the essential features of the theory formulated here. For simplicity, in the Fig. 2A-B only Respiratory Complex I is considered, which oxidizes NADH by initiating the Electron Transport Chain (ETC), coupled with the transfer of 4 H⁺ for each molecule of NADH oxidized. On Fig. 2A (scheme of canonical chemiosmotic theory with coupling with FoF1-ATP synthase, protons are transferred to the intermembrane space by the action of Complex I; then the return of protons to the matrix, rotating the Fo Subunit of the FoF1-ATP Synthase nanomachine which synthesizes ATP in the matrix. This coupling called traditionally *delocalized coupling* cannot be accepted because no accumulation of protons is possible, otherwise the mitochondrion would collapse due to the high local acidity. Even Robert Williams, who was one of the most ardent opponents of the chemiosmotic theory, insisted that protons could not accumulate in the intermembrane space [40][41]. His famous objection is that protons transferred into intermembrane space would be lost in a "Pacific Ocean" [42]. In confirmation it has been highlighted that in the microvolume of a mitochondrion the protons can be very few [43].

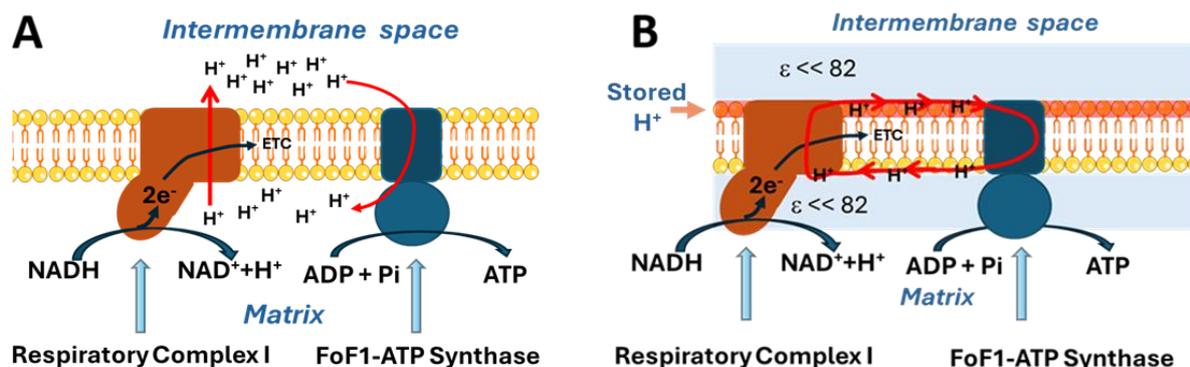


Figure 2. Only Respiratory complex I is considered, which oxidizes NADH by initiating the Electron Transport Chain (ETC), coupled with the transfer of 4 H⁺ for each molecule of NADH oxidized. In Fig. 2A the protons accumulated in the intermembrane space - which would generate the proton motive force – and return to the matrix, rotating the Fo subunit which transmits a molecular constriction to the F1 subunit with the consequent synthesis of ATP. This coupling called "delocalized coupling" should be rejected because no accumulation of free protons is possible, otherwise the mitochondrion would collapse due to the too low pH. In Fig. 2B, the existence on both sides of the membrane of a layer with dielectric constant $\epsilon \ll 82$ is taken into account, confirming the insulation of the membrane with respect to the aqueous bulk. The protons moved by Respiratory Complex I migrate along phosphate and nitrogen

121 ionizable residues (such as ethanolamine and choline) of phospholipids in the intermembrane-
122 facing hemilayer, pass through the Fo subunit of ATP synthase—with ATP synthesis by the
123 F1 subunit—and return to ionizable residues of phospholipids in the matrix-facing hemilayer.
124 But such groups can retain protons in the ionizable residues of phospholipids (highlight in
125 orange) and this supports their function as proton capacitors. This scheme is consistent with
126 “localized coupling”.

127
128 In Fig. 2B (scheme of “Proton Capacitor Theory”) the protons moving from
129 Complex I travel along the ionizable head of the phospholipid facing the
130 intermembrane space without being released into it, which is a well documented
131 process [44][45][46][47]. The protons are captured by the Fo subunit of the FoF1-ATP
132 synthase nanomachine, which synthesizes ATP, and can return to Respiratory
133 Complex I by traveling along the ionizable “head”, i.e. phosphate and basic group as
134 ethanolamine and choline, of the phospholipid facing the matrix. This groups can retain
135 proton by accumulating them and this is the core of the Proton Capacitor Theory. It is
136 important to take into account that the inner mitochondrial membrane is isolated from
137 the aqueous phase by a layer of ~ 7–10 nm with a dielectric constant $\epsilon < 82$ [48][49].
138 The “proton motive force” exerted by Respiratory Complex I is difficult to formulate in
139 thermodynamic/mathematical terms. This coupling can be traced back to *localized*
140 *coupling* which has already been proposed [50][51][52][53] as an alternative to
141 *delocalized coupling* and is based on the rigid separation of the processes that occur
142 in the membrane from those that occur in the aqueous phase.
143

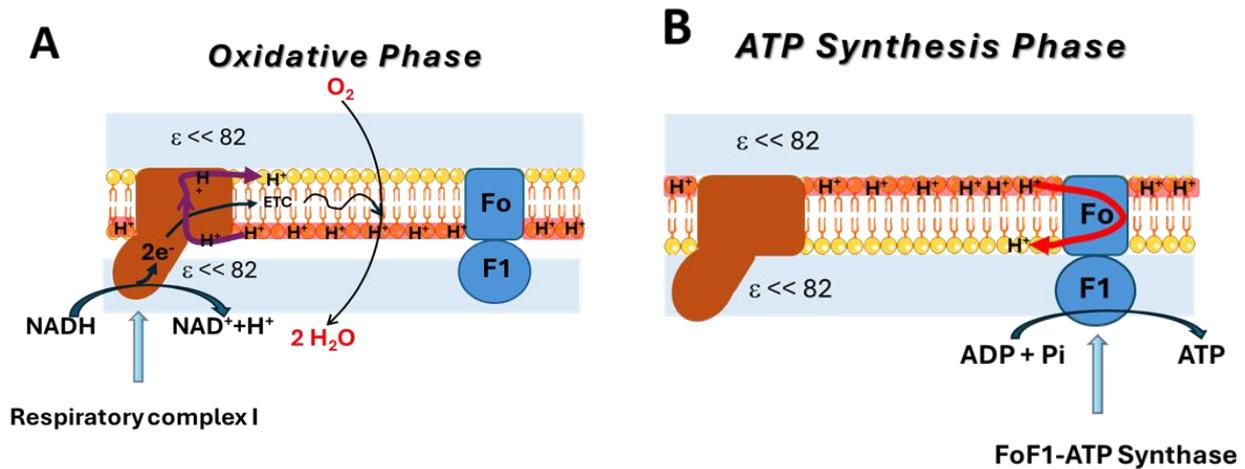
144 4. Two phases of OXPHOS and Basics of Membrane’s Proton 145 Capacitor 146

147 Having established that the proton movement characteristic of OXPHOS is
148 confined exclusively to the membrane and that the protons move thanks to the polar
149 residues of the phospholipids which are located on the sides of the membrane bilayer,
150 it is intuitive that these groups can express a buffer function [19], that is, they can retain
151 protons in one phase to release them in a subsequent phase which leads to the rotation
152 of the Fo subunit of the ATP Synthase with consequent ATP synthesis.

153 This function of membrane is called “Proton Capacitor” and to configure it’s
154 function is necessary to assume that OXPHOS can occur in two phases, as has been
155 proposed for sleep [12] and generally for the course of OXPHOS so much so that an
156 intermediate energy accumulator and its division into two phases has already been
157 proposed [54].
158

159 Figure 3A shows the scheme of the Oxidative Phase, with proton translocation
160 (performed by Respiratory Complex I) from the polar groups of the phospholipids
161 facing the matrix to the polar groups of the phospholipids facing the intermembrane
162 space. Figure 3B reports the scheme of the ATP Synthesis phase, with the protons
163 passing from the polar “heads” of the phospholipids, facing to intermembrane space,
164 to the polar “heads” of the phospholipids facing the matrix. The ATP Synthesis phase

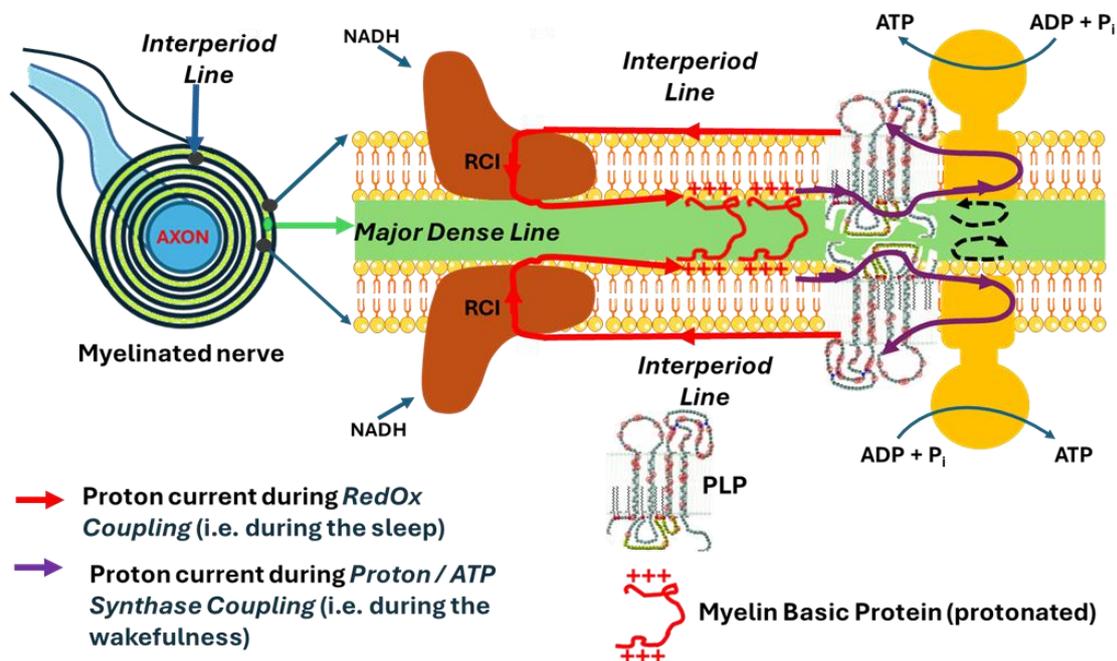
165 is operative in the presence of ADP + Pi in the matrix, while in their absence the
 166 OXPPOS is limited to the Oxidative Phase with accumulation of protons in the
 167 dissociable groups of the membranes.
 168



169 **Figure 3. The diagram of the phases that lead to the "charging" and "discharging" of**
 170 **the proton capacitor is shown.** For simplicity, only Respiratory Complex I is considered,
 171 which oxidizes NADH by initiating the Electron Transport Chain (ETC), coupled with the
 172 transfer of 4 H⁺ for each molecule of NADH oxidized. Fig. 3A provides the scheme of the
 173 "Oxidative Phase": the protons accumulated in the ionic group of phospholipid hemi-layer
 174 facing the matrix are transferred to the ionic group of phospholipid hemi-layer facing the
 175 intermembrane space and there they accumulate. Fig. 3B provides the scheme of the "ATP
 176 Synthesis Phase": the protons accumulated in the ionic group of phospholipid hemi-layer
 177 facing the intermembrane space are transferred to the ionic group of phospholipid in the hemi-
 178 layer facing the matrix passing through the Fo subunit with consequent it's rotation and
 179 transmission a molecular constriction to the F1 subunit that synthesizes ATP in the matrix.

180
 181
 182 The capacitance of the proton capacitor can be increased by the contribution of
 183 membrane proteins that can function as proton buffers as they contain numerous
 184 dissociable acidic groups [55]. This is the case of myelin which contains high amounts
 185 of Myelin Basic Protein (MBP) and proteolipid (PLP) which provide a notable increase
 186 in the proton capacitance of the myelin sheath, as has been roughly calculated [12].
 187 Figure 4 shows the functioning of the proton capacitor that supports the sleep/wake
 188 cycle: the sleep phase corresponds to the oxidative phase, already described here in
 189 mitochondria, and is characterized by a proton pathway represented by the red line,
 190 while the wake phase, corresponding to the mitochondrial ATP synthesis phase, is
 191 represented by the purple line.

192



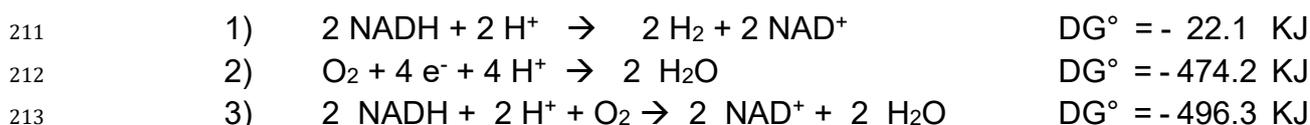
193 **Figure 4. Scheme of proton capacitor function in the myelin sheath to support the**
 194 **sleep/wake cycle.** The myelin sheath surrounding the nerve is shown on the left and a detail
 195 of the arrangement of the membranes is shown on the right: the interperiod line is the area
 196 corresponding to the mitochondrial matrix, while the major dense line corresponds to the
 197 mitochondrial intermembrane space. **Red line** = “proton recharging” phase of myelin: protons
 198 are transferred by means of Respiratory Complex I from the basic domain of the PLP immersed
 199 in the interperiod line to the MBP and to the basic domain of the PLP, both immersed in the
 200 major dense line of myelin. **Purple line** = Myelin “proton discharge” phase with coupled ATP
 201 synthesis: protons are transferred from the MBP and the basic domain of the PLP immersed
 202 in the major dense line to the basic domain of the PLP immersed in the interperiod line;
 203 the proton current flows through the Fo subunit of the ATP Synthase with the consequent ATP
 204 synthesis by the F1 subunit immersed in the interperiod line.

205

206 5. Thermodynamic evaluations

207

208 Referring to the schemes presented above, we evaluate the thermodynamics
 209 of the entire process. The reactions with their corresponding free energy variations are
 210 as follows:



214 Equation 1) refers to the oxidation of 2 moles of NADH, which sets in motion 4
 215 electrons that travel along the ETC until they discharge onto the O₂ molecule,
 216 producing 2 molecules of H₂O.

217 Equation 2) refers to the subsequent discharge of 4 electrons onto the O₂
218 molecule, producing 2 molecules of H₂O. The sum (3) cancels out 4 electrons + 4 H⁺
219 from equation 2) with 2 H₂ from equation 1). Conventionally, one refers to one reduced
220 oxygen atom producing one molecule of H₂O, so the DG° in equation 3) is divided by
221 2. Therefore, the DG° relating to one oxygen atom is DG° = - 496.3 : 2 = - 248.2 kJ.
222 The protons set in motion for 1 NADH oxidized are 4 from Respiratory Complex I
223 (NADH dehydrogenase), 3 from Respiratory Complex III (cytochrome c
224 oxidoreductase) and 4 from Respiratory Complex IV (cytochrome c oxidase) for a total
225 of 11. The stoichiometry of protons transferred by the Respiratory Complexes just cited
226 refers to an authoritative contribution [56].

227 For many species, the complete rotation of Fo requires 8 protons, as many as
228 the c subunits that make up Fo [57], therefore 11 protons ensure 11/8 = 1.38 rotations
229 of the Fo subunit. Each rotation produces 3 ATP from the F1 subunit, therefore with 11
230 protons we will have the synthesis of 1.38 x 3 = 4.12 ATP. Each ATP involves a DG°
231 = 30.5 KJ therefore the energy recovered in the form of ATP is 30.5 x 4.12 = 125.7 KJ.
232 A theoretical efficiency of $h = 125.7 : 248.2 = 51\%$ is obtained which is within those
233 predicted with other methods [58].

234

235 6. Conclusions and outlook

236

237 The traditional statement "*The Mitochondria are the Powerhouse of the cells*" can
238 be amplified into "*The Mitochondria are exporters to the cell of the Power Molecular
239 Machinery*" because it assemble the numerous proteins that carry out Oxidative
240 Phosphorylation (OXPHOS) which through vesiculation they transfer to other cellular
241 and extracellular sites making them able to synthesize ATP on their own. Furthermore,
242 it appears that extra-mitochondrial ATP synthesis is widespread both within the cell
243 and in the body's circulation due to the diverse presence of MDV.

244 All this requires overcoming the chemiosmotic theory [38][59][60][61] [62]
245 [63][64] [65] and so the Proton Capacitor theory has now been formulated which
246 foresees a clear division between the processes that occur in the membrane and those
247 that occur in the aqueous phase that laps both sides of the inner mitochondrial
248 membrane or other cellular membranes where OXPHOS is operative. The extra-
249 mitochondrial OXPHOS have been found in the plasma membrane [66] [67] [68] [69]
250 [70] [71] [72], in the endoplasmic reticulum [73][74], in the sarcoplasmic reticulum [11]
251 and in the widely spread MDVs [29][75][28][16].

252 It goes without saying that this new approach greatly simplifies the original
253 chemiosmotic theory because i) the processes involved occur only in the proteolipidic
254 phase of the membrane and ii) it offers a new interpretation of the variation of the
255 fluorescent signal generated by the so-called "membrane potential indicators" which
256 with good reason reflect the degree of protonation of the membrane itself, as has been
257 pointed out [19].

258 The import-export process of the OXPHOS molecular machinery has recently
259 been outlined in the article “*Mitochondria-derived vesicles: potential nano-batteries to*
260 *recharge the cellular powerhouse*” [17] and introduces the concept of “batteries” which
261 is reminiscent of the possible existence of the “Proton Capacitor” which has already
262 been described as the fulcrum around which the accumulation of energy in the form of
263 protons revolves which are incorporated into the acid/base buffers of the membrane
264 whose function has been proposed to formulate a coherent theory of sleep [12].
265 Furthermore the existence of the proton capacitor in membranes also depends on the
266 formation of inverted micelles which increase the proton storage capacity [63].

267 The capacity of membranes to store protons, and therefore energy, is greater the
268 larger the developed surface area of the membrane and this appears to be an
269 evolutionary advantage accumulated by mitochondria which have developed, unlike
270 the bacteria from which they derive, an internal membrane folded into cristae with the
271 consequent increase in surface area for the same volume.

272 Interestingly, the sleep hypothesis requires a theory of OXPHOS functioning that
273 coincides with the proton capacitor theory illustrated here, and this result may pave
274 the way for understanding energy homeostasis in those energy-intensive organs, such
275 as the heart, skeletal muscle, and kidneys, where extramitochondrial OXPHOS is
276 highly active.

277 278 **Acknowledgements**

279 The Author are indebted to Prof. Ann Saada (University of Gerusalem, Israel)
282 and Richard Funk (University of Dresden, Germany) for his support and insightful
283 discussions.

284 285 **References**

- 286
287 [1] Y. Watanabe, E. Tajika, Atmospheric oxygenation of the early earth and earth-like
288 planets driven by competition between land and seafloor weathering, in: *Earth, Planets*
289 *Sp.*, Springer, 2021: p. 188.
- 290 [2] L. Margulis, Symbiotic theory of the origin of eukaryotic organelles; criteria for proof.,
291 *Symp. Soc. Exp. Biol.* (1975) 21–38. <http://www.ncbi.nlm.nih.gov/pubmed/822529>.
- 292 [3] N. Lane, *Power, sex, suicide: Mitochondria and the meaning of Life*, 2nd ed., Oxford
293 Landmark Science, Oxford, 2018.
- 294 [4] A. Morelli, C. Rosano, A True Symbiosis for the Mitochondria Evolution, *Bioenerg. Open*
295 *Access* 05 (2016). <https://doi.org/10.4172/2167-7662.1000137>.
- 296 [5] T.J.G. Ettema, Evolution: Mitochondria in the second act., *Nature* 531 (2016) 39–40.
297 <https://doi.org/10.1038/nature16876>.
- 298 [6] A.A. Pittis, T. Gabaldón, Late acquisition of mitochondria by a host with chimaeric
299 prokaryotic ancestry., *Nature* 531 (2016) 101–4. <https://doi.org/10.1038/nature16941>.
- 300 [7] I. Panfoli, D. Calzia, P. Bianchini, S. Ravera, A. Diaspro, G. Candiano, A. Bachi, M.
301 Monticone, M.G. Aluigi, S. Barabino, G. Calabria, M. Rolando, C. Tacchetti, A. Morelli,

- 302 I.M. Pepe, Evidence for aerobic metabolism in retinal rod outer segment disks., *Int. J.*
303 *Biochem. Cell Biol.* 41 (2009) 2555–65. <https://doi.org/10.1016/j.biocel.2009.08.013>.
- 304 [8] S. Ravera, I. Panfoli, D. Calzia, M.G. Aluigi, P. Bianchini, A. Diaspro, G. Mancardi, A.
305 Morelli, Evidence for aerobic ATP synthesis in isolated myelin vesicles, *Int J Biochem*
306 *Cell Biol* 41 (2009) 1581–1591. [https://doi.org/S1357-2725\(09\)00012-0](https://doi.org/S1357-2725(09)00012-0)
307 [pii]10.1016/j.biocel.2009.01.009.
- 308 [9] S. Ravera, M.G. Signorello, M. Bartolucci, S. Ferrando, L. Manni, F. Caicci, D. Calzia,
309 I. Panfoli, A. Morelli, G. Leoncini, Extramitochondrial energy production in platelets, *Biol.*
310 *Cell* 110 (2018) 97–108. <https://doi.org/10.1111/boc.201700025>.
- 311 [10] M. Ishikawa, Z.S. Hasanali, Y. Zhao, A. Das, M. Lavaert, C.J. Roman, J. Londregan, D.
312 Allman, A. Bhandoola, Bone marrow plasma cells require P2RX4 to sense extracellular
313 ATP, *Nature* 626 (2024) 1102–1107. <https://doi.org/10.1038/s41586-024-07047-2>.
- 314 [11] H. Lee, S.-H. Kim, J.-S. Lee, Y.-H. Yang, J.-M. Nam, B.-W. Kim, Y.-G. Ko, Mitochondrial
315 oxidative phosphorylation complexes exist in the sarcolemma of skeletal muscle., *BMB*
316 *Rep.* 49 (2016) 116–21. <https://doi.org/10.5483/bmbrep.2016.49.2.232>.
- 317 [12] A.M. Morelli, A. Saada, F. Scholkmann, Myelin: A possible proton capacitor for energy
318 storage during sleep and energy supply during wakefulness., *Prog. Biophys. Mol. Biol.*
319 (2025). <https://doi.org/10.1016/j.pbiomolbio.2025.03.001>.
- 320 [13] J. Heyn, M.A. Heuschkel, C. Goettsch, Mitochondrial-Derived Vesicles-Link to
321 Extracellular Vesicles and Implications in Cardiovascular Disease., *Int. J. Mol. Sci.* 24
322 (2023). <https://doi.org/10.3390/ijms24032637>.
- 323 [14] L.-C. Tábara, M. Segawa, J. Prudent, Molecular mechanisms of mitochondrial
324 dynamics., *Nat. Rev. Mol. Cell Biol.* 26 (2025) 123–146. <https://doi.org/10.1038/s41580-024-00785-1>.
- 326 [15] J. Gao, L. Wang, J. Liu, F. Xie, B. Su, X. Wang, Abnormalities of Mitochondrial Dynamics
327 in Neurodegenerative Diseases, *Antioxidants* 6 (2017) 25.
328 <https://doi.org/10.3390/antiox6020025>.
- 329 [16] A. Sugiura, G.-L. McLelland, E.A. Fon, H.M. McBride, A new pathway for mitochondrial
330 quality control: mitochondrial-derived vesicles., *EMBO J.* 33 (2014) 2142–56.
331 <https://doi.org/10.15252/embj.201488104>.
- 332 [17] S. Mishra, G. Deep, Mitochondria-derived vesicles: potential nano-batteries to recharge
333 the cellular powerhouse, *Extracell. Vesicles Circ. Nucleic Acids* 5 (2024) 271–275.
334 <https://doi.org/10.20517/evcna.2023.71>.
- 335 [18] T. König, H.M. McBride, Mitochondrial-derived vesicles in metabolism, disease, and
336 aging., *Cell Metab.* 36 (2024) 21–35. <https://doi.org/10.1016/j.cmet.2023.11.014>.
- 337 [19] A.M. Morelli, A. Saada, F. Scholkmann, Extra-mitochondrial ATP synthesis, proton
338 dynamics at the membrane, and mitochondria-derived vesicles: Current findings and
339 considerations, *Mitochondrial Commun.* (2025).
340 <https://doi.org/10.1016/j.mitoco.2025.06.002>.
- 341 [20] R. Iorio, S. Petricca, G. Di Emidio, S. Falone, C. Tatone, Mitochondrial Extracellular
342 Vesicles (mitoEVs): Emerging mediators of cell-to-cell communication in health, aging
343 and age-related diseases., *Ageing Res. Rev.* 101 (2024) 102522.
344 <https://doi.org/10.1016/j.arr.2024.102522>.
- 345 [21] L. de Meis, L.A. Ketzer, R.M. da Costa, I.R. de Andrade, M. Benchimol, Fusion of the
346 Endoplasmic Reticulum and Mitochondrial Outer Membrane in Rats Brown Adipose
347 Tissue: Activation of Thermogenesis by Ca²⁺, *PLoS One* 5 (2010) e9439.
348 <https://doi.org/10.1371/journal.pone.0009439>.

- 349 [22] T.-J. Bae, M.-S. Kim, J.-W. Kim, B.-W. Kim, H.-J. Choo, J.-W. Lee, K.-B. Kim, C.S. Lee,
350 J.-H. Kim, S.Y. Chang, C.-Y. Kang, S.-W. Lee, Y.-G. Ko, Lipid raft proteome reveals
351 ATP synthase complex in the cell surface, *Proteomics* 4 (2004) 3536–3548.
352 <https://doi.org/10.1002/pmic.200400952>.
- 353 [23] A.G. Lee, Myelin: Delivery by raft, *Curr Biol* 11 (2001) R60-2.
- 354 [24] M. Vinson, O. Rausch, P.R. Maycox, R.K. Prinjha, D. Chapman, R. Morrow, A.J. Harper,
355 C. Dingwall, F.S. Walsh, S.A. Burbidge, D.R. Riddell, Lipid rafts mediate the interaction
356 between myelin-associated glycoprotein (MAG) on myelin and MAG-receptors on
357 neurons., *Mol. Cell. Neurosci.* 22 (2003) 344–52. [https://doi.org/10.1016/s1044-
358 7431\(02\)00031-3](https://doi.org/10.1016/s1044-7431(02)00031-3).
- 359 [25] M. Colombini, Structure and mode of action of a voltage dependent anion-selective
360 channel (VDAC) located in the outer mitochondrial membrane., *Ann. N. Y. Acad. Sci.*
361 341 (1980) 552–63. <https://doi.org/10.1111/j.1749-6632.1980.tb47198.x>.
- 362 [26] S. Hiller, J. Abramson, C. Mannella, G. Wagner, K. Zeth, The 3D structures of VDAC
363 represent a native conformation., *Trends Biochem. Sci.* 35 (2010) 514–21.
364 <https://doi.org/10.1016/j.tibs.2010.03.005>.
- 365 [27] V. Shoshan-Barmatz, V. De Pinto, M. Zweckstetter, Z. Raviv, N. Keinan, N. Arbel,
366 VDAC, a multi-functional mitochondrial protein regulating cell life and death., *Mol.*
367 *Aspects Med.* 31 (2010) 227–85. <https://doi.org/10.1016/j.mam.2010.03.002>.
- 368 [28] P. D’Acunzo, R. Pérez-González, Y. Kim, T. Hargash, C. Miller, M.J. Alldred, H.
369 Erdjument-Bromage, S.C. Penikalapati, M. Pawlik, M. Saito, M. Saito, S.D. Ginsberg,
370 T.A. Neubert, C.N. Goulbourne, E. Levy, Mitovesicles are a novel population of
371 extracellular vesicles of mitochondrial origin altered in Down syndrome., *Sci. Adv.* 7
372 (2021). <https://doi.org/10.1126/sciadv.abe5085>.
- 373 [29] R. Hazan Ben-Menachem, D. Lintzer, T. Ziv, K. Das, I. Rosenhek-Goldian, Z. Porat, H.
374 Ben Ami Pilo, S. Karniely, A. Saada, N. Regev-Rudzki, O. Pines, Mitochondrial-derived
375 vesicles retain membrane potential and contain a functional ATP synthase., *EMBO Rep.*
376 24 (2023) e56114. <https://doi.org/10.15252/embr.202256114>.
- 377 [30] P.J. Yao, C. Noguerras-Ortiz, K.A. Pucha, D. Kapogiannis, ATP Synthase Abundance in
378 Neuronal Extracellular Vesicles Reflects Changes in the Mitochondria of Parent
379 Neurons., *J. Extracell. Vesicles* 14 (2025) e70140. <https://doi.org/10.1002/jev2.70140>.
- 380 [31] P. Mitchell, Coupling of phosphorylation to electron and hydrogen transfer by a chemi-
381 osmotic type of mechanism, *Nature* 191 (1961) 144–148.
- 382 [32] A.M. Morelli, S. Ravera, D. Calzia, I. Panfoli, Hypothesis of lipid-phase-continuity proton
383 transfer for aerobic ATP synthesis, *J Cereb Blood Flow Metab* 33 (2013) 1838–1842.
- 384 [33] A.M. Morelli, S. Ravera, D. Calzia, I. Panfoli, An update of the chemiosmotic theory as
385 suggested by possible proton currents inside the coupling membrane, *Open Biol.* 9
386 (2019) 180221. <https://doi.org/10.1098/rsob.180221>.
- 387 [34] A.M. Morelli, S. Ravera, I. Panfoli, The aerobic mitochondrial ATP synthesis from a
388 comprehensive point of view., *Open Biol.* 10 (2020) 200224.
389 <https://doi.org/10.1098/rsob.200224>.
- 390 [35] S. Nath, Energy landscapes and dynamics of ion translocation through membrane
391 transporters: a meeting ground for physics, chemistry, and biology, *J. Biol. Phys.* 47
392 (2021) 401–433. <https://doi.org/10.1007/s10867-021-09591-8>.
- 393 [36] A.M. Morelli, M. Chiantore, S. Ravera, F. Scholkmann, I. Panfoli, Myelin sheath and
394 cyanobacterial thylakoids as concentric multilamellar structures with similar bioenergetic
395 properties., *Open Biol.* 11 (2021) 210177. <https://doi.org/10.1098/rsob.210177>.

- 396 [37] V. Wray, Field guide to Nath's research work on ATP synthesis and hydrolysis.,
397 Biosystems. 252 (2025) 105461. <https://doi.org/10.1016/j.biosystems.2025.105461>.
- 398 [38] S. Nath, Rethinking the Classical Chemiosmotic Theory, Biol. Theory (2025).
399 <https://doi.org/10.1007/s13752-025-00499-3>.
- 400 [39] S. Ji, Chemiosmotic vs conformational models of oxidative phosphorylation: Theory and
401 mechanistic insights., Biosystems. 259 (2026) 105637.
402 <https://doi.org/10.1016/j.biosystems.2025.105637>.
- 403 [40] R.J. Williams, Proton-driven phosphorylation reactions in mitochondrial and chloroplast
404 membranes., FEBS Lett. 53 (1975) 123–5.
- 405 [41] R.J. Williams, The multifarious couplings of energy transduction., Biochim. Biophys.
406 Acta 505 (1978) 1–44.
- 407 [42] R.J. Williams, The multifarious couplings of energy transduction., Biochim. Biophys.
408 Acta 505 (1978) 1–44. [https://doi.org/10.1016/0304-4173\(78\)90007-1](https://doi.org/10.1016/0304-4173(78)90007-1).
- 409 [43] W. Bal, E. Kurowska, W. Maret, The Final Frontier of pH and the Undiscovered Country
410 Beyond, PLoS One 7 (2012) e45832. <https://doi.org/10.1371/journal.pone.0045832>.
- 411 [44] S. Serowy, S.M. Saparov, Y.N. Antonenko, W. Kozlovsky, V. Hagen, P. Pohl, Structural
412 proton diffusion along lipid bilayers., Biophys. J. 84 (2003) 1031–7.
413 [https://doi.org/10.1016/S0006-3495\(03\)74919-4](https://doi.org/10.1016/S0006-3495(03)74919-4).
- 414 [45] T.H. Nguyen, C. Zhang, E. Weichselbaum, D.G. Knyazev, P. Pohl, P. Carloni, Interfacial
415 water molecules at biological membranes: Structural features and role for lateral proton
416 diffusion., PLoS One 13 (2018) e0193454.
417 <https://doi.org/10.1371/journal.pone.0193454>.
- 418 [46] S.J. Ferguson, Chemiosmotic coupling. Protons fast and slow., Curr. Biol. 5 (1995) 25–
419 7. [https://doi.org/10.1016/s0960-9822\(95\)00008-x](https://doi.org/10.1016/s0960-9822(95)00008-x).
- 420 [47] J. Teissié, M. Prats, P. Soucaille, J.F. Tocanne, Evidence for conduction of protons
421 along the interface between water and a polar lipid monolayer., Proc. Natl. Acad. Sci.
422 U. S. A. 82 (1985) 3217–21. <https://doi.org/10.1073/pnas.82.10.3217>.
- 423 [48] D.A. Cherepanov, B.A. Feniouk, W. Junge, A.Y. Mulkidjanian, Low dielectric permittivity
424 of water at the membrane interface: effect on the energy coupling mechanism in
425 biological membranes., Biophys. J. 85 (2003) 1307–16. [https://doi.org/10.1016/S0006-3495\(03\)74565-2](https://doi.org/10.1016/S0006-3495(03)74565-2).
- 427 [49] A.Y. Mulkidjanian, D.A. Cherepanov, Probing biological interfaces by tracing proton
428 passage across them., Photochem. Photobiol. Sci. 5 (2006) 577–87.
429 <https://doi.org/10.1039/b516443e>.
- 430 [50] B. Rieger, W. Junge, K.B. Busch, Lateral pH gradient between OXPHOS complex IV
431 and F(0)F(1) ATP-synthase in folded mitochondrial membranes., Nat. Commun. 5
432 (2014) 3103. <https://doi.org/10.1038/ncomms4103>.
- 433 [51] S.A. Eremeev, L.S. Yaguzhinsky, On Local Coupling of Electron Transport and ATP-
434 Synthesis System in Mitochondria. Theory and Experiment., Biochemistry. (Mosc). 80
435 (2015) 576–81. <https://doi.org/10.1134/S0006297915050089>.
- 436 [52] S. Nath, Coupling mechanisms in ATP synthesis: Rejoinder to “Response to molecular-
437 level understanding of biological energy coupling and transduction,” Biophys. Chem.
438 272 (2021) 106579. <https://doi.org/10.1016/j.bpc.2021.106579>.
- 439 [53] J. Sjöholm, J. Bergstrand, T. Nilsson, R. Šachl, C. von Ballmoos, J. Widengren, P.
440 Brzezinski, The lateral distance between a proton pump and ATP synthase determines
441 the ATP-synthesis rate., Sci. Rep. 7 (2017) 2926. [12](https://doi.org/10.1038/s41598-017-</p></div><div data-bbox=)

- 442 02836-4.
- 443 [54] D.B. Kell, A protet-based, protonic charge transfer model of energy coupling in oxidative
444 and photosynthetic phosphorylation., *Adv. Microb. Physiol.* 78 (2021) 1–177.
445 <https://doi.org/10.1016/bs.ampbs.2021.01.001>.
- 446 [55] P.L. Jordan, H.N. Raum, S. Gröger, U. Weininger, Protonation Kinetics in Proteins at
447 Basic pH Determined by pH-Dependent NMR Relaxation Reveal the Entire Relationship
448 between Kinetics and pK a Values., *JACS Au* 5 (2025) 2334–2341.
449 <https://doi.org/10.1021/jacsau.5c00245>.
- 450 [56] M. Wikström, G. Hummer, Stoichiometry of proton translocation by respiratory complex
451 I and its mechanistic implications, *Proc. Natl. Acad. Sci.* 109 (2012) 4431–4436.
452 <https://doi.org/10.1073/pnas.1120949109>.
- 453 [57] I.N. Watt, M.G. Montgomery, M.J. Runswick, A.G.W. Leslie, J.E. Walker, Bioenergetic
454 cost of making an adenosine triphosphate molecule in animal mitochondria., *Proc. Natl.*
455 *Acad. Sci. U. S. A.* 107 (2010) 16823–7. <https://doi.org/10.1073/pnas.1011099107>.
- 456 [58] D.F. Wilson, Oxidative phosphorylation: regulation and role in cellular and tissue
457 metabolism, *J. Physiol.* 595 (2017) 7023–7038. <https://doi.org/10.1113/JP273839>.
- 458 [59] E. Bertero, C. Maack, Rethinking Mitchell's Chemiosmotic Theory: Potassium
459 Dominates Over Proton Flux to Drive Mitochondrial F1Fo-ATP Synthase, *Function* 3
460 (2022). <https://doi.org/10.1093/function/zqac012>.
- 461 [60] G. Drochioiu, Reinterpretation of Jagendorf's classic experiment on
462 photophosphorylation, *BioSystems* 257 (2025) 105614.
463 <https://doi.org/10.1016/j.biosystems.2025.105614>.
- 464 [61] R.M. Farahani, An Addendum to the Chemiosmotic Theory of Mitochondrial Activity:
465 The Role of RNA as a Proton Sink, *Biomolecules* 15 (2025) 87.
466 <https://doi.org/10.3390/biom15010087>.
- 467 [62] E.S. Gasanoff, "Acidic and Basic Proteins from Spider *Latrodectus Pallidus* Venom that
468 Induce Formation of Non-Bilayer Phase and Increase Proton Capacity in Model Myelin
469 Membranes Feature High Sequence Homology to Isoforms of Myelin Basic Protein:
470 Pharmacological Relevan, *Biomed. J. Sci. Tech. Res.* 56 (2024).
471 <https://doi.org/10.26717/BJSTR.2024.56.008895>.
- 472 [63] E.S. Gasanoff, R.K. Dagda, Cobra Venom Cytotoxins as a Tool for Probing Mechanisms
473 of Mitochondrial Energetics and Understanding Mitochondrial Membrane Structure,
474 *Toxins (Basel)*. 16 (2024) 287. <https://doi.org/10.3390/toxins16070287>.
- 475 [64] Z. Zeng, M. Wei, S. Zhang, H. Cui, R.K. Dagda, E.S. Gasanoff, Bee Venom Proteins
476 Enhance Proton Absorption by Membranes Composed of Phospholipids of the Myelin
477 Sheath and Endoplasmic Reticulum: Pharmacological Relevance., *Pharmaceuticals*
478 *(Basel)*. 18 (2025). <https://doi.org/10.3390/ph18091334>.
- 479 [65] E.A. Kasumov, R.E. Kasumov, I. V. Kasumova, A mechano-chemiosmotic model for the
480 coupling of electron and proton transfer to ATP synthesis in energy-transforming
481 membranes: a personal perspective, *Photosynth. Res.* 123 (2015) 1–22.
482 <https://doi.org/10.1007/s11120-014-0043-3>.
- 483 [66] N. Arakaki, T. Nagao, R. Niki, A. Toyofuku, H. Tanaka, Y. Kuramoto, Y. Emoto, H.
484 Shibata, K. Magota, T. Higuti, Possible role of cell surface H⁺ -ATP synthase in the
485 extracellular ATP synthesis and proliferation of human umbilical vein endothelial cells.,
486 *Mol. Cancer Res.* 1 (2003) 931–9. <http://www.ncbi.nlm.nih.gov/pubmed/14638865>.
- 487 [67] B.-W. Kim, H.-J. Choo, J.-W. Lee, J.-H. Kim, Y.-G. Ko, Extracellular ATP is generated
488 by ATP synthase complex in adipocyte lipid rafts., *Exp. Mol. Med.* 36 (2004) 476–85.

- 489 <https://doi.org/10.1038/emm.2004.60>.
- 490 [68] E. Champagne, L. Martinez, X. Collet, R. Barbaras, Ecto-F1Fo ATP synthase/F1
491 ATPase: metabolic and immunological functions., *Curr Opin Lipidol.* 17 (2006) 279–84.
492 <https://doi.org/10.1097/01.mol.0000226120.27931.76>.
- 493 [69] R. Mangiullo, A. Gnoni, A. Leone, G. V Gnoni, S. Papa, F. Zanotti, Structural and
494 functional characterization of F(o)F(1)-ATP synthase on the extracellular surface of rat
495 hepatocytes, *Biochim Biophys Acta* 1777 (2008) 1326–1335.
- 496 [70] B.-W. Kim, C.S. Lee, J.-S. Yi, J.-H. Lee, J.-W. Lee, H.-J. Choo, S.-Y. Jung, M.-S. Kim,
497 S.-W. Lee, M.-S. Lee, G. Yoon, Y.-G. Ko, Lipid raft proteome reveals that oxidative
498 phosphorylation system is associated with the plasma membrane., *Expert Rev.*
499 *Proteomics* 7 (2010) 849–66. <https://doi.org/10.1586/epr.10.87>.
- 500 [71] S. Ravera, M.G. Aluigi, D. Calzia, P. Ramoino, A. Morelli, I. Panfoli, Evidence for ectopic
501 aerobic ATP production on c6 glioma cell plasma membrane, *Cell Mol Neurobiol* 31
502 (2011) 313–321.
- 503 [72] Y.-W. Chang, T. Tony Yang, M.-C. Chen, Y. Liaw, C.-F. Yin, X.-Q. Lin-Yan, T.-Y. Huang,
504 J.-T. Hou, Y.-H. Hung, C.-L. Hsu, H.-C. Huang, H.-F. Juan, Spatial and temporal
505 dynamics of ATP synthase from mitochondria toward the cell surface, *Commun. Biol.* 6
506 (2023) 427. <https://doi.org/10.1038/s42003-023-04785-3>.
- 507 [73] H. Ishii, S. Kunihiro, M. Tanaka, K. Hatano, T. Nishikata, Cytosolic subunits of ATP
508 synthase are localized to the cortical endoplasmic reticulum-rich domain of the ascidian
509 egg myoplasm., *Dev Growth Differ* 54 (2012) 753–66.
510 <https://doi.org/10.1111/dgd.12003>.
- 511 [74] K. Allen-Worthington, J. Xie, J.L. Brown, A.M. Edmunson, A. Dowling, A.M. Navratil, K.
512 Scavelli, H. Yoon, D.-G. Kim, M.S. Bynoe, I. Clarke, M.S. Roberson, The F0F1 ATP
513 Synthase Complex Localizes to Membrane Rafts in Gonadotrope Cells, *Mol.*
514 *Endocrinol.* 30 (2016) 996–1011. <https://doi.org/10.1210/me.2015-1324>.
- 515 [75] R. Hazan (Ben-Menachem), O. Pines, A. Saada, Mitochondrial derived vesicles- Quo
516 Vadis?, *FEBS J.* 291 (2024) 4660–4669. <https://doi.org/10.1111/febs.17103>.

517

518 **Copyright** ©2026 The author. This Open Access communication (not peer-
519 reviewed) is distributed under the terms of the Creative Commons
520 Attribution License, which permits unrestricted use, distribution, and
521 reproduction in any medium, provided the original authors and source are
522 credited. © remains with the authors, who have granted BEC preprints an
523 Open Access publication license in perpetuity.