

REVIEW ARTICLE Multivariate meta-analyses of mitochondrial complex I and IV in major depressive disorder, bipolar disorder, schizophrenia, Alzheimer disease, and Parkinson disease

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Complex I (NADH dehydrogenase, NDU) and complex IV (cytochrome-c-oxidase, COX) of the mitochondrial electron transport chain have been implicated in the pathophysiology of major psychiatric disorders, such as major depressive disorder (MDD), bipolar disorder (BD), and schizophrenia (SZ), as well as in neurodegenerative disorders, such as Alzheimer disease (AD) and Parkinson disease (PD). We conducted meta-analyses comparing complex I and IV in each disorder MDD, BD, SZ, AD, and PD, as well as in normal aging. The electronic databases Pubmed, EMBASE, CENTRAL, and Google Scholar, were searched for studies published between 1980 and 2018. Of 2049 screened studies, 125 articles were eligible for the meta-analyses. Complex I and IV were assessed in peripheral blood, muscle biopsy, or postmortem brain at the level of enzyme activity or subunits. Separate meta-analyses of mood disorder studies, MDD and BD, revealed moderate effect sizes for similar abnormality patterns in the expression of complex I with SZ in frontal cortex, cerebellum and striatum, whereas evidence for complex IV alterations was low. By contrast, the neurodegenerative disorders, AD and PD, showed strong effect sizes for shared deficits in complex I and IV, such as in peripheral blood, frontal cortex, cerebellum, and substantia nigra. Beyond the diseased state, there was an age-related robust decline in both complexes I and IV. In summary, the strongest support for a role for complex I and/or IV deficits, is in the pathophysiology of PD and AD, and evidence is less robust for MDD, BD, or SZ.

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INTRODUCTION

Mitochondrial dysfunction is implicated in the pathophysiology of major psychiatric disorders, such as major depressive disorder (MDD) [1], bipolar disorder (BD) [2] and schizophrenia (SZ) [3], as well as neurodegenerative disorders, such as Alzheimer disease (AD) [4] and Parkinson disease (PD) [4]. Mitochondria are intracellular organelles that produce adenosine triphosphate (ATP), the main source of cellular energy. Impaired mitochondrial function results in decreased ATP production, impaired bioenergetics, apoptosis and oxidative stress [5]. Prior to the generation of ATP, the electrons extracted from nutrients are transported along the electron transport chain (ETC) and the energy released is directed into a transmembrane proton.

Research has identified two enzymes of the ETC located at the inner mitochondrial membrane as being particular impaired in these five disorders MDD, BD, SZ, AD, and PD. The first enzyme, complex I (NADH dehydrogenase, NDU) consists of 45 subunits, seven of which are encoded by mitochondrial DNA (mtDNA) and the remaining subunits by nuclear DNA (nDNA). Complex I is one of the entry enzymes of cellular respiration or oxidative phosphorylation in the mitochondrion. It is also the largest multimeric enzyme complex of the ETC and is a major contributor to the generation of the proton gradient across the mitochondrial inner membrane, which drives ATP production. The second enzyme of interest, complex IV (cytochrome-c-oxidase, COX)

consists of 13 subunits, three of which are encoded by mtDNA, the remainder by nDNA. Complex IV catalyzes the final step in the mitochondrial ETC and, due to its rate-limiting role in this oxidative process [6], has been proposed as a key markers of mitochondrial function [7]. Numerous excellent reviews [1–4] have discussed the details of impairments in both complex I and IV enzyme activities and subunit assembly within each of the above mentioned disorders. A meta-analysis summarizing the findings across these disorders could not be found in the literature. The remaining complexes II (succinate dehydrogenase), III (cyto-chrome c reductase) and V (ATP synthase) either have not been studied, or they have been studied to a much smaller degree in these five disorders, compared with complex I and IV, and thus there are too few data for a meta-analysis.

We chose these five disorders MDD, BD, SZ, AD, and PD not only because of the potential common mitochondrial dysfunction, but also based on their clinical similarities. Though regarded as different disorders in major classification systems like DSM and ICD, there is also overlap in clinical symptoms. Depression is found in mood disorders, but is also frequent in schizophrenia and both AD and PD [8, 9]. Psychotic symptoms are observed in MDD and BD, as well as SZ ((DSM-IV-TR), [10]). Although, Alzheimer's and Parkinson's diseases have distinct brain histopathology, both are age-related neurodegenerative conditions characterized by memory loss and depression and have some commonality in molecular

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pathogenesis, such as proteinopathy, neuroinflammation, and oxidative stress [11]. Cross-disorder commonalities are also found in schizophrenia sharing dopaminergic abnormalities with Parkinson's [12] and cognitive impairment (dementia praecox) with Alzheimer's [13].

Beyond disease states, normal aging plays an independent role in mitochondrial ETC function [14]. To separate the influence of age from that of disease, we therefore additionally meta-analyzed complex I and IV functioning in healthy older versus younger individuals. The present meta-analyses aimed to provide a synthesis of the work done on complex I and IV that can guide future research and stimulate development of novel in vivo technologies such as for example the assessment of the redox states of complex I NAD+/NADH ratio [15, 16] or oxidized complex IV [17].

MATERIALS AND METHODS

Literature search and study identification

We conducted a structured literature search in the PubMed, EMBASE, CENTRAL, and Google Scholar, to identify studies published between January 1980 and January 2018 using the search strings 'NADH dehydrogenase' OR 'cytochrome-c-oxidase' OR 'complex I' OR 'complex IV' AND 'major depressive disorder' OR 'bipolar disorder' OR 'schizophrenia' OR 'Alzheimer' OR 'Parkinson' OR 'age/aging'. We manually reviewed reference lists in all retrieved articles for related publications.

Inclusion criteria were: studies investigated either complex I and/ or complex IV, studies published in the English language, studies reporting original human data and studies that investigated patients in comparison with a control group. Exclusion criteria were: animal studies, cellular studies, genetic studies, studies with less than three subjects per group, case reports, letters to the editor and editorials and publications not reporting original data.

Data extraction

Aggregated data were extracted for each of the disorders MDD, BD, SZ, AD, PD and normal aging (AGING). Aggregated data contained quantitative data (i.e., number of patients and controls, mean age of patients, and controls), qualitative information (i.e., methods such as enzyme activity or subunits at the level of mRNA or protein expression, and tissues assayed, such as peripheral blood, muscle, or brain regions of interest, ROIs) and outcomes (effect sizes in terms of the standardized mean difference, SMD, and *p*-values). Enzyme activity was extracted as a corrected ratio to citrate synthase to normalize for mitochondrial content, if available. For studies on AGING, we chose a breakpoint of mean 60 years [18] to derive group differences between older (> 60) and younger (< 60) individuals using t-test. In case of medication reported in the studies, both unmedicated and medicated cohorts were included. In case of BD, data were included from both BD I and BD II during depressive episodes, but not manic episodes.

The authors of some studies were contacted for permission to reuse original data [19–25]; no answer was received in some cases and the data could therefore not or only partially included [19, 20, 23, 25]. Some data were read from figures [18, 26–45].

Multivariate random-effect meta-analyses

Separate meta-analyses were first computed for each disorder. Moderator variables were generated representing the combination of two aspects: (1) method (enzyme activity or subunit); and (2) tissue (blood, muscle, or ROI). A moderator was included in the meta-analysis only if it was reported by at least three studies. This threshold was chosen after extensive prior testing in order to avoid spurious results.

A summary meta-analysis was computed across all disorders. Here, the moderator variable consisted of the five disorders MDD, BD, SZ, AD, PD, and AGING. This analysis included all data found to be eligible in the present work and thus also included rare outcomes that could not be included in the separate disease/ aging meta-analyses.

For all meta-analyses, a multivariate random-effect model was applied based on the Metafor package [46] as implemented in R [47]. The multivariate model accounts for heterogeneity and dependency in the underlying true effects of multiple moderators that can overlap within subjects and studies (random factors). To adjust for age effects, the age-difference between patients and controls was added as a continuous covariate. To adjust for sample size, the effects were weighted based on study size. To allow for heterogeneity differences between moderators, an unstructured variance-covariance matrix was applied (rma.mv function; observed effects = SMD + age-difference; variance-covariance matrix = COV; weight = study size; variance structure = UN; mods = moderators/ disorder; random factors = subject/study; method = REML).

Forest plots were generated to illustrate the results of the multivariate models in terms of the weighted beta coefficients with 95% confidence intervals (CIs) and *p*-values. Heterogeneity was assessed using Cochran's *Q*-test and the inconsistency l^2 statistic that directly indicates to what extent each outcome contributes to the total variance. Publication bias was assessed using Egger's regression analysis.

RESULTS

Data extraction

Of 2049 screened studies, 125 studies were eligible for the metaanalyses, some of them for more than one disorder (MDD N = 7, BD N = 19, SZ N = 21, AD N = 43, PD N = 35, AGING N = 18) (Table 1). The overall number of studies reporting on complex I (total *N* = 101, MDD *N* = 6, BD *N* = 19, SZ *N* = 15, AD *N* = 19, PD *N* = 32, AGING N = 11) and complex IV (total N = 104, MDD N = 4, BD *N* = 10, SZ *N* = 10, AD *N* = 39, PD *N* = 28, AGING *N* = 15) were similar but varied between disorders. Both complex I and IV were analysed in either peripheral blood elements, such as platelets, lymphocytes, or other blood cells (MDD N = 1, BD N = 10, SZ N =8, AD N = 9, PD N = 14, AGING N = 2), in muscle biopsies (MDD N = 2, BD N = 0, SZ N = 0, AD N = 0, PD N = 9, AGING N = 14), or in postmortem brain samples (MDD N = 4, BD N = 8, SZ N = 13, AD N = 34, PD N = 14, AGING N = 2). Both complex I and IV were analysed at the level of enzymatic activity (either in blood, muscle, or brain, MDD N = 4, BD N = 5, SZ N = 11, AD N = 32, PD N = 32, AGING N = 14), or of their subunits by mRNA level (only brain, MDD *N* = 3, BD *N* = 7, SZ *N* = 7, AD *N* = 7, PD *N* = 1, AGING *N* = 2) or protein level (only brain, MDD N = 3, BD N = 12, SZ N = 7, AD N= 6, PD N = 3, AGING N = 5). In the following, enzyme activity is denoted with NDU/COX respectively, whereas subunits as assessed by either mRNA or protein expression are referred to with their full names.

Note that we combined data from levels of mRNA/protein expression for all subunits. The correlation between mRNA and protein expression is generally thought to be around 40% [48, 49], which is typically attributed to other levels of regulation between transcript and protein product [50]. We separated the findings in a pre-analysis but found no significant differences between mRNA and protein expression and therefore presented the results together.

Assay of enzyme activity was more common in studies of neurodegenerative disorders, AD (88%) and PD (66%); whereas, the assessment of mRNA/protein expression was more common in studies of mood disorders, MDD (60%), BD (79%) and SZ (56%). None of the five disorders, however, showed differences in enzyme activity or mRNA/protein expression for complex I or IV (p > 0.05). Moreover, there were no differences between peripheral blood and brain in the enzyme activity/mRNA/protein levels of complex I and IV (p > 0.05).

All studies involved subjects over 18 years old, except two studies examined early-onset SZ patients below age 18 years [27, 51]. The mean age-difference between cases and controls differed

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Table 1. Eligible studies						
MDD (<i>N</i> = 7)	BD (<i>N</i> = 19)	SZ (N = 21)	AD (N = 43)	PD (<i>N</i> = 35)	Aging (<i>N</i> = 18)	
Andreazza [45]	Akarsu [71]	Altar [23]	Aksenov [112]	Anderson [131]	Boffoli [156]	
Ben-Shachar [170]	Altar [23]	Andreazza [45]	Alikhani [109]	Barroso [138]	Boffoli [153]	
Ben-Shachar [59]	Andreazza [45]	Andreazza [26]	Armand-Ugon [115]	Benecke [118]	Cabré [18]	
Gardner [171]	Andreazza [26]	Ben-Shachar [170]	Bosetti [106]	Bindoff [125]	Distefano [39]	
Gardner [172]	Beech [76]	Ben-Shachar [74]	Cardoso [96]	Blake [120]	Emelyanova [154]	
Karry [24]	Ben-Shachar [170]	Ben-Shachar [59]	Casademont [173]	Blin [126]	Gueugneau [41]	
Sanchez-Bahillo [174]	Ben-Shachar [59]	Cavelier [103]	Cavelier [103]	Bravi [121]	Hsieh [42]	
	de Sousa [175])	Dror [75]	Chagnon [36]	Bronstein [129]	Lanza [155]	
	Gubert [176]	Gubert [176]	Chandrasekaran [113]	Cardellach [127]	Merlo Pich [177]	
	lwamoto [60]	lwamoto [60]	Chandrasekaran [37]	Chagnon [36]	Merlo Pich [178]	
	Karry [24]	Karry [24]	Cooper [54]	Cooper [179]	Ogborn [149]	
	Kim [43]	Kim [43]	Cottrell [114]	DiDonato [55]	Ojaimi [30]	
	Konradi [25]	Maurer [180]	Devi [38]	Duke [181]	Pestronk [150]	
	Munkholm [28]	Mehler-Wex [27]	Feldhaus [97]	Gatt [40]	Rasmussen [182]	
	Naydenov [183]	Prince [67]	Fišar [98]	Gu [110]	Rooyackers [151]	
	Rosenfeld [73]	Prince [68]	Fukuyama [184]	Haas [133]	Safdar [32]	
	Sun [58]	Rice [185]	Gu [110]	Hanagasi [134]	Trounce [35]	
	Washizuka [22]	Rosenfeld [73]	Hirai [56]	Keeney [139]	Zucchini [152]	
	Washizuka [21]	Taurines [51]	Kim [186]	Krige [122]		
		Washizuka [21]	Kish [44]	Mann [130]		
		Whatley [187]	Kish [108]	Martín [128]		
			Liang [188]	Mizuno [135]		
			Long [104]	Mythri [29]		
			Lunnon [57]	Nakagawa-Hattori [123]		
			Mancuso [91]	Parker [116]		
			Manczak [19]	Parker [31]		
			Maurer [99]	Schaegger [136]		
			Mutisya [107]	Schapira 1990a [140]		
			Parker [92]	Schapira 1990b [137]		
			Parker 1994a [93]	Shinde [132]		
			Parker 1994b [100]	Shoffner [124]		
			Pérez-Gracia [101]	Thomas [20]		
			Reichmann [52]	Varghese [117]		
			Sekar [189]	Wiedemann [53]		
			Sheehan [94]	Yoshino [119]		
			Simonian [33]			
			Simonian [34]			
			Terni [190]			
			Valla [105]			
			Valla [95]			
			Verwer [111]			
			Vitali [191]			
			Wong-Riley [102]			

more than one disorder. The detailed data extracted from these studies are provided in the supplementary materials (Tables S1-6).

between disorders (main effect F = 35.21, p < 0.0001). Psychiatric studies, MDD (mean age-difference \pm STD = -0.7 ± 3.9), BD (mean age-difference \pm STD = 0.8 ± 4.9) and SZ (mean age-difference \pm STD = -1.0 ± 3.7) included closer age-matched cases and controls compared with AD (mean age-difference \pm STD = 5.0 ± 7.5) or PD (mean age-difference \pm STD = 5.3 ± 5.9) as indicated by significant post hoc comparisons (p < 0.0001). Some studies did not provide

mean ages for cases and/or controls [52–56], for which we therefore assumed zero age-difference.

Multivariate random-effect meta-analyses

Meta-analyses were performed for each of the disorders MDD, BD, SZ, AD, PD, and AGING. Based on the precondition that a moderator was reported at least three times across studies, the

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Fig. 1 Forest plots MDD, BD, and SZ. Multivariate random-effect estimates of the SMD (95% CI, *p*-values) of complex I (NDU) and IV (COX) enzyme activity and subunits are shown both in numerical and graphical form. Values lower than 0 indicate that patients had lower levels than controls (CO), and vice versa for values greater than 0; the dashed vertical line at SMD = 0 indicates no effect. The size of the filled circles for each estimated SMD is proportional to the weight of the studies. *P* putamen, *NA* nucleus accumbens, *GP* globus pallidus

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Fig. 2 Forest plots AD, PD, and AGING. Multivariate random-effect estimates of the SMD (95% CI, *p*-values) of complex I (NDU) and IV (COX) enzyme activity and subunits are shown both in numerical and graphical form. Values lower than 0 indicate that patients (or older (O) individuals) had lower levels than controls (CO) (or younger (Y) individuals), and vice versa for values greater than 0; the dashed vertical line at SMD = 0 indicates no effect. The size of the filled circles for each estimated SMD is proportional to the weight of the studies

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number of studies finally included in the meta-analyses (and the percentage of the eligible studies) differed between disorders MDD (complex I N = 4 (67%), complex IV N = 2 (50%)), BD (complex I N = 16 (84%), complex IV N = 3 (30%)), SZ (complex I N = 14 (93%), complex IV N = 6 (60%)), AD (complex I N = 12 (63%), complex IV N = 33 (85%)), PD (complex I N = 29 (91%), complex IV N = 24 (86%), and AGING (complex I N = 7 (64%), complex IV N = 14 (93%)).

Mood disorders. In MDD (Fig. 1a), complex I subunits NDUFS1_{cer}ebellum (p = 0.0075), NDUFV1cerebellum (p = 0.00014), and NDUFV2_{cerebellum} (p = 0.00051) were lower in cerebellum and subunit NDUFV1frontal (p = 0.0047) was lower in frontal cortex compared with controls. No differences were found for complex IV (note that only in this case, we set the moderator threshold to at least two).

In BD (Fig. 1b), complex I subunit NDUFS1_{cerebellum} (p < 0.0001) was lower in cerebellum, subunits NDUFS1_{frontal} (p = 0.033), and NDUFS7_{frontal} (p < 0.0001) were lower in frontal cortex and subunit NDUFS1_{striatum} (p = 0.0023) was lower in striatum compared with controls. No differences were found for complex IV.

In SZ (Fig. 1c), complex I subunit NDUFV2_{blood} (p = 0.019) was lower in blood, whereas subunit NDUFS1_{blood} (p = 0.00086) was higher in blood compared with controls. Further, subunits NDUFS1_{striatum} (p = 0.00072), NDUFV1_{striatum} (p < 0.0001), and NDUFV2_{striatum} (p < 0.0001) were lower in striatum, subunit NDUFV1_{frontal} (p = 0.011) was lower in frontal cortex, whereas subunit NDUFV2_{parietal} (p = 0.019) was higher in parietal cortex compared with controls. Complex IV enzyme activity was lower in

frontal cortex (COX_{frontal} p = 0.016), but higher in basal ganglia (COX_{NA/GP/P} p < 0.0001, nucleus accumbens, globus pallidus and putamen) compared with controls.

We also assessed the overlap in findings between MDD, BD, and SZ using meta-regressions. Results confirmed lower NDUFS1/ NDUFV1/NDUFV2 levels in the cerebellum in MDD (p = 0.005/p = 0.001/p = 0.006) and in some subunits also BD (p = 0.088/p = 0.001/p = 0.258) compared to SZ, respectively. By contrast, lower NDUFS1/ NDUFV1/NDUFV2 levels in striatum were observed in SZ compared to MDD (p = 0.001/p < 0.0001/p < 0.0001) and in some subunits also BD (p = 0.250/p < 0.0001/p < 0.0001), respectively. Further, both BD (p = 0.024) and SZ (p = 0.008) revealed higher levels of NDUFV2 in the parietal cortex compared to MDD. Last, in BD, subunit NDUFS1_{blood} levels (p = 0.002) in blood and subunit NDUFS7_{frontal} (p = 0.009) in frontal cortex were lower compared to SZ.

Neurodegenerative disorders. In AD (Fig. 2a), complex I enzyme activity in the temporal cortex (NDU_{temporal/entorhinal} p = 0.0023) and subunits NDUFA1/4/7-9_{blood} (p < 0.0001), NDUFB2/3/6_{blood} (p < 0.0001) and NDUFS3/4/5_{blood} (p < 0.0001) were lower in blood compared with controls. It should be noted that these subunits were reported by Lunnon et al. [57], which was the largest study eligible for the present work with a total of 209 subjects (patients N = 104, controls N = 105). Initially only a small subset of that study could be included in the meta-analysis because of the high moderator threshold; in order to include more of the results from that study, we summarized results across subunits to build sufficient moderators.



Fig. 3 Forest plot Summary. Multivariate random-effect estimates of the SMD (95% CI, *p*-values) of complex I and IV are shown both in numerical and graphical form. SMD are ordered according to their strengths. Values lower than 0 indicate that patients (P) had lower levels than controls (CO), and vice versa for values greater than 0; the dashed vertical line at SMD = 0 indicates no effect. The size of the filled circles for each estimated SMD is proportional to the weight of the studies

Table 2. Heterogeneity and Publication bias.					
		Q	<i>p</i> -value	l ² (%)	
Heteroge	neity and inconsister	псу			
MDD	Complex I	549.160	0.000	91%	
	Complex IV	0.030	1.000	0%	
BD	Complex I	137,410.000	0.000	100%	
	Complex IV	873.150	0.000	95%	
sz	Complex I	2197.500	0.000	97%	
	Complex IV	209.560	0.000	88%	
AD	Complex I	64,812.000	0.000	100%	
	Complex IV	4490.300	0.000	97%	
PD	Complex I	437.650	0.000	80%	
	Complex IV	3083.300	0.000	99%	
AGING	Complex I	27.934	0.218	18%	
	Complex IV	11.684	1.000	0%	
Egger's re	egression				
		z-value	df	<i>p</i> -value	
MDD	Complex I	-1.969	4	0.120	
	Complex IV	0.630	2	0.593	
BD	Complex I	-2.191	17	0.043	
	Complex IV	-0.042	8	0.968	
SZ	Complex I	0.135	13	0.895	
	Complex IV	1.035	8	0.331	
AD	Complex I	-0.109	17	0.915	
	Complex IV	-1.381	37	0.176	
PD	Complex I	-2.127	30	0.042	
	Complex IV	-0.176	26	0.862	
AGING	Complex I	-1.025	9	0.332	
	Complex IV	-2.820	13	0.014	

(Top) Heterogeneity and Inconsistency. Significant *Q* statistics indicate the existence of heterogeneity A l^2 -value of 0% indicates no observed inconsistency; whereas larger values shows increasing inconsistency. (Bottom) Egger's regression test for . The existence of potential publication bias is indicated by *p*-values <0.05 *df* degrees of freedom

Complex IV enzyme activity was lower in frontal cortex (COX_{frontal} p = 0.044), motor cortex (COX_{motor} p = 0.0003), occipital cortex (COX_{occiptal} p = 0.032), parietal cortex (COX_{parietal} p = 0.00069), temporal/entorhinal cortex (COX_{temporal/entorhinal} p = 0.00012), and hippocampus (COX_{hippocampus} p < 0.0001) compared to controls. Subunit COX2_{hippocampus} in hippocampus (p < 0.0001) and subunit COX7_{blood} in peripheral blood (p = 0.011) were lower.

In PD (Fig. 2b), complex I enzyme activity in peripheral muscle (NDU_{muscle} p = 0.000084) as well as in substantia nigra (NDU_{sub-}stantia nigra p = 0.006) was lower compared to controls. Complex IV enzyme activity in frontal cortex (COXfrontal p = 0.036) was also lower.

The overlap between AD and PD was assessed using metaregressions. Results showed comparable patterns of enzyme activity in blood (COX_{blood} p = 0.981) and brain such as in frontal cortex (NDU_{frontal} p = 0.171, COX_{frontal} p = 0.711), cerebellum (COX_{cerebellum} p = 0.582) and substantia nigra (COX_{substantia} nigra p = 0.671), indicating no significant differences between AD and PD.

Aging. To assess complex I and IV functioning in normal aging, we performed a separate meta-analysis in healthy subjects using a breakpoint of 60 years [18] (Fig. 2c). Older (\geq 60 years) subjects in comparison with younger (<60 years) subjects had lower complex

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I enzyme activity in frontal cortex (NDU_{frontal} p < 0.0001) and muscle (NDU_{muscle} p < 0.0001, NDUFB8_{muscle} p = 0.00015). Similarly, complex IV enzyme activity was lower in frontal cortex (COX_{frontal} p = 0.00072) and muscle (COX_{muscle} p = 0.00027, COX2_{muscle} p < 0.0001) in older compared to younger subjects.

Summary comparison of all cases and controls. A summary metaanalysis was performed to categorize the disorders according to the severity of overall mitochondrial impairments (Fig. 3). Robust deficits were observed for complex I in PD (p = 0.0016) and AGING (p = 0.009) and for complex IV in AD (p = 0.004) and AGING (p < 0.0001). MDD (p < 0.0001) also showed an effect on complex I; however, it was based on small samples sizes from few studies and can therefore not be considered a robust result. The main effect across both complex I and IV was significant in PD (p = 0.00058), AD (p = 0.049) and AGING (p < 0.0001) indicating that these three conditions were also affected if both complex I and IV are considered.

Heterogeneity and publication bias

Overall, there was a high degree of heterogeneity (Table 2). The l^2 for the five disorders MDD, BD, SZ, AD, and PD ranged between 80 and 100%. Noticeable, there was low heterogeneity in AGING in both complex I (Q = 27.934, p = 0.218, $l^2 = 18\%$) and IV (Q = 11.684, p = 1, $l^2 = 0\%$). Putative low heterogeneity was observed in MDD for complex IV (Q = 0.030, p = 1, $l^2 = 0\%$), most likely explained by the fact that the data came from few studies (N = 4).

Publication bias as assessed using Egger's regression test (Table 2) provided non-significant results for most of the data indicating no publication bias. Considerable publication bias was only found in PD for complex I (p = 0.043) and in AGING for complex IV (p = 0.014).

DISCUSSION

Our findings provide a picture of the current findings on complex I and IV in mood and neurodegenerative disorders (Fig. 4). We found that mood disorders, MDD, and BD, share commonalities with SZ in having low expression of complex I subunits. Neurodegenerative disorders, AD, and PD, share similarities in

MDD	Small number of studies with small sample sizes Region-specific heterogeneity Moderate effects in complex I, small effects in complex IV
BPD	 Small sample sizes Region-specific heterogeneity Moderate effects in complex I, small effects in complex IV
SZ	 Small sample sizes Region-specific heterogeneity Moderate effects in complex I and IV
AD	 Large number of studies with large sample sizes Tissue-specific heterogeneity (low in blood, high in brain) Strong effects in complex I and IV
PD	Large number of studies No heterogeneity Strong effects in complex I and IV
AGING	Most robust effects No heterogeneity Strongest effects in complex I and IV

Fig. 4 Schematic overview of our findings. Highlighted are the main findings of the meta-analyses for the five disorders MDD, BD, SZ, AD, PD, and normal aging. Region-specific heterogeneity means that complex levels vary from brain region to brain region (i.e., variation from ROI to ROI); tissue-specific heterogeneity indicates that complex levels vary between tissues (i.e., peripheral blood versus brain)

complex I and IV enzyme activity abnormalities. Due to heterogeneity in study findings, strong evidence for complex I and/or IV deficits is present for AD and PD; whereas, the overall proof for the involvement of these complexes in MDD, BD, and SZ is less robust and requires further research. Normal aging is a substantial independent factor contributing to both complex I and IV decline and should be considered when interpreting study heterogeneity.

Mood disorders

Mood disorders show shared patterns of complex I subunits NDUFS1, NDUFV1 and NDUFV2 between MDD and BD on the one hand, and between BD and SZ on the other hand (Fig. 1) [24, 26, 43, 45, 58-601. While, MDD and to some degree BD, have lower levels of NDUFS1, NDUFV1, NDUFV2 in cerebellum compared with SZ, the latter had lower NDUFS1, NDUFV1, NDUFV2 levels in striatum compared with MDD and to some degree BD; this indicates a shared mechanism between MDD and BD that is distinct from SZ. By contrast, BD and SZ had elevated levels of NDUFV2 in parietal cortex compared with MDD, which may indicate a shared mechanism between BD and SZ. These brain regions, i.e., the frontal cortex [61, 62], cerebellum [63, 64], and striatum [65], have been previously suggested to contribute to the pathophysiology of mood disorders and schizophrenia spectrum [66]. Distinct from the mood disorders, SZ has higher complex IV levels in basal ganglia [67, 68] perhaps related to previous neuropathological findings in putamen, nucleus accumbens, and globus pallidus [69, 70].

Notably, while most of the eligible studies pointed to dysregulation of complex I and/or IV, the reported direction of change has not always been consistent. While some of the heterogeneity may be due to potential differences in the clinical picture like presence of psychosis in MDD or BD and presence of depression in SZ, several other explanations may underlie these discrepancies. First, there may be state-related differences in the expression of complex I and IV. For example BD patients showed up-regulation of complex I subunits during manic compared to depressive episodes [71] maybe related to elevated brain metabolism in frontal and temporal cortex [72]. Similarly in SZ, state-dependence has been suggested with increased complex I activity in active psychotic patients but decreased activity in patients with residual schizophrenia [73-75]. Unfortunately, postmortem studies were unable to provide information on whether patients were depressed, manic or euthymic at time of death. Second, there may be brain region-specific heterogeneity, possibly indicating that ETC energy production varies from region to region, as previously exemplarily shown [59]. Third, there may be tissue-specific differences between blood samples from living patients and brain samples obtained postmortem. For example, it has been hypothesized that ETC genes are upregulated in blood, but downregulated in some or all brain regions in BD [76] and SZ [77], although this was not confirmed statistically by the present meta-analysis. Fourth, the majority of patients were receiving psychotropic medication, such as antidepressants, mood stabilizers, and antipsychotics. All of these pharmacological agents have been reported to potentially interfere or even inhibit the mitochondrial ETC [78]. Finally, differences in brain pH at the time of death may explain difference in gene expression [79]. Larger sample studies of medication-free patients are needed to clarify more precisely the relationships between diagnosis, mood state, treatment status, and brain pH in mood disorders and schizophrenia pathophysiology. There is also evidence of shared global gene expression patterns between mood disorders and schizophrenia that go beyond purely mitochondrial function that could be investigated in relation [80].

Neurodegenerative disorders

Neurodegenerative disorders AD and PD show similarities in the impairment of complex I and IV enzyme activities in the blood, frontal

cortex, cerebellum, and substantia nigra (Fig. 2). These brain regions are potentially involved in the cognitive decline and neuronal loss-in-AD [81, 82], as well as in the motor and non-motor related degeneration of neurotransmitter systems in PD [83, 84].

In AD, there are a well-documented deficit in both complex I and IV [4, 85, 86]. These deficits are thought to be due to the neuronal toxicity induced by amyloid β peptide (A β), an important component of AD pathogenesis, in that AB accumulates in mitochondria, directly inhibits mitochondrial enzymes, perpetuates oxidative stress, and leads to a hypometabolic state, which causes mitochondrial dysfunction [87-90]. The meta-analysis supported these findings in terms of downregulated complex I and IV in the blood [57, 91–98], frontal cortex [38, 44, 56, 99–104], motor cortex [102, 105, 106], occipital cortex [44, 107, 108], parietal cortex [36, 44, 52, 102, 107, 108], temporal/entorhinal . cortex [37, 44, 52, 54, 99, 102, 107–110] and hippocampus [33, 34, 36, 44, 52, 99, 102, 106, 111–114]. The strength of the effect sizes are in line with findings of neuronal loss in entorhinal cortex, hippocampus and association neocortex [81]. Due to limited data, the meta-analysis was not able to distinguish between Brack stages in AD. A large study in AD (N = 148) reported less expression of complex I and IV subunits in entorhinal and frontal cortex in disease stages V–VI but not in stages I–II [115]. A smaller study (N = 18) compared complex I and IV enzyme activity in frontal cortex in stages III-VI to stages I-II but found no stagedependence [19]. There were also no differences between earlyversus late-onset AD (age <60 versus age >60) [96].

In PD, the meta-analysis points to downregulation of both complex I and IV enzyme activity in the blood, muscle, and brain. For example, inhibition of complex I has been reported in platelets [110, 116-122] and skeletal muscles [123-127], although there was dissenting studies in platelets [128, 129] or muscle [55, 130, 131]. Similarly, complex IV has been found to be decreased in blood [116, 118, 121, 122, 132, 133] and muscle [124, 125, 127, 131], while others could not confirm this for either platelets [120, 134] or muscle [53, 55, 123, 126, 130]. Within the brain, some studies reported lower complex I enzyme activity in substantia nigra [110, 130, 135–137], as well as low activity and impaired assembly of complex I in frontal cortex [29, 31, 40, 135, 138, 139], the latter not being confirmed in the meta-analysis. Similarly, complex IV in the brain is reported to be low in frontal cortex [31, 36, 135, 138, 139] and substantia nigra [36, 110, 130, 136, 140], the latter finding not confirmed here. Note that although all data eligible for the meta-analysis were obtained from patients suffering from the more common sporadic form of PD, derangements in mitochondrial metabolism and oxidative stress have also been strongly linked to the less common familial form of PD [141].

Likewise for mood disorders, medication may affect mitochondrial functioning in neurodegeneration. Cholinesterase (ChE) inhibitors and amyloid-beta binding alcohol dehydrogenase (ABAD) modulators are used to treat the key symptoms in AD in order to improve activities of daily living, behavior, and cognition [142]. Most of these drugs can negatively affect both complex I and IV [143]. There are however newly developed ChEs and ABADs, as well as other more recent drug developments that target multiple AD pathophysiological pathways that are thought to have a better outcome on mitochondrial function [144]. By contrast, in PD, levodopa, cardidopa, and selegiline, which are used to treat PD motor symptoms, have not been shown not affect complex I and IV in PD [145, 146]. Similarly, melatonin that is used as sleep/wake regulator in PD, acts as an effective antioxidant and mitochondrial function protector [147]. These observations support the hypothesis that impaired complex I and IV activity in PD patients is a characteristic of the disease and not due to medications.

Aging

Normal aging has been associated with a decline in mitochondrial quality [14] and activity of both complex I and IV [6, 148]. The

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meta-analysis supported these findings in terms of decreased levels of enzyme activity in the muscle [32, 39, 41, 42, 149–155] and frontal cortex [18, 30, 153, 156] (Fig. 2c). Across all metaanalyses, the effect of aging was the most robust both in terms of the strength of the summary effect sizes (Fig. 3), as well as the low degree of heterogeneity (Table 2).

The decline in both complex I and IV in normal aging [6, 148] as indicated by this meta-analysis can represent a functionally serious impairment of mitochondrial function with respect to basal ATP production [157]. Therefore, aged neurons with lower mitochondrial mass and enzyme activities as well as more dysfunctional mitochondria may be unable to respond adequately to any increased ATP demand [158].

The term normal aging should be considered carefully. Normal aging is thought to be a function of both chronological age as well as biological age [32]. Although the underlying mechanism is still controversial, age-dependent mitochondrial decline may contribute to loss-of-physical capacity and mental exertion [159, 160]. The 'free radical' theories of aging [161, 162] state that reactive oxygen species (ROS) produced aberrantly during mitochondrial electron transport damage mitochondrial components, including mtDNA. Since, mtDNA encodes essential complex I and IV proteins [163], its damage and mutagenesis would disrupt ETC complex assembly and, in turn, lead to more ROS production, cellular oxidative stress and tissue dysfunction that promote aging [164]. This theory, however, has been thought to be too simplistic and there is debate about other age-dependent cellular changes that may be important in precipitating the decline in mitochondrial ETC [160].

Summary comparison of all cases and controls

Summarizing the overall severity of mitochondrial impairment in the five disorders (Fig. 3) indicates that major psychiatric disorders MDD, BD, and SZ show less robust effects. The small effect sizes in those pathologies were most likely due to the small number of studies with small sample sizes, large region-specific heterogeneity and variation in medication and treatment, which made in harder to detect underlying mitochondrial disease. By contrast, strong evidence for a complex I deficit is found in PD, whereas a complex IV deficit is present in AD; with the simultaneous decline in complex I and IV being significant in both AD and PD. These results indicate an interaction of complex I with IV involved in the assembly and stability of complex I [165–167].

The interpretation of the strengths of the effect sizes between the five disorder compared with normal aging, as shown in Fig. 3, should however be cautious. Aging is a major risk factor for developing both AD [168] and idiopathic PD [169]. Consequently, the effects of normal aging and that of the diseases cannot be clearly separated because of their inter-dependence. Of note, patients in all studies included were age-matched, thus in principle presenting age-adjusted effects. We therefore argue that the lesser effects in mood disorders is not related to the younger age of these patients, and the larger effects in AD is also not related to older age.

Further research is required to enlarge sample sizes and strengthen the evidence base. Our results may stimulate the application of novel in vivo technologies such as assessing complex I NAD+/NADH redox state via auto-fluorescence [16] or magnetic resonance spectroscopy [15] and complex IV redox states via functional near-infrared spectroscopy [17].

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ADDITIONAL INFORMATION

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