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Blood eosinophils and chronic obstructive pulmonary disease: a GOLD Science Committee 2022 review

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Complete List of Authors:	Singh, Dave; The University of Manchester Agusti, Alvar; Fundacio Clinic per a la Recerca Biomedica, Martinez, Fernando J.; Cornell Medical College, Papi, Alberto; University of Ferrara, Research Centre on Asthma and COPD Pavord, Ian; Oxford University, Nuffield department of Medicine, Respiratory Medicine Wedzicha, Jadwiga; Imperial College London, National Heart and Lung Institute Vogelmeier, Claus; University of Marburg, Pulmonary Diseases Halpin, David; University of Exeter College of Medicine, University of Exeter Medical School; Royal Devon and Exeter Hospital,
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4	Dave Singh ¹ , Alvar Agusti ² , Fernando J. Martinez ³ , Alberto Papi ⁴ , Ian D Pavord ⁵ , Jadwiga A
5	Wedzicha ⁶ , Claus F. Vogelmeier ⁷ , David MG Halpin ⁸
6	Affiliations
7	1. University of Manchester, Manchester University NHS Foundation Trust, Manchester, UK
8	2. Respiratory Institute, Hospital Clinic, University of Barcelona, IDIBAPS, CIBERES, Spain
9	3. Weill Cornell Medicine, NY Presbyterian Hospital, New York, NY, USA
10	4. Respiratory Medicine Unit, University of Ferrara, University Hospital S.Anna, Ferrara, Italy
11	5. Oxford Respiratory NIHR BRC and Nuffield Department of Medicine, University of Oxford.
12	6. National Heart and Lung Institute, Imperial College, London, UK
13 14	 Department of Medicine, Pulmonary and Critical Care Medicine, University of Marburg, Member of the German Center for Lung Research (DZL), Marburg, Germany
15 16	8. University of Exeter Medical School, College of Medicine and Health, University of Exeter, Exeter, UK
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26

27 Abstract

COPD is a heterogeneous condition. Some patients benefit from treatment with inhaled corticosteroids (ICS) but this requires a precision medicine approach, based on clinical characteristics (phenotyping) and biological information (endotyping) in order to select patients most likely to benefit. The GOLD 2019 report recommended using exacerbation history combined with blood eosinophil counts (BEC) to identify such patients. Importantly, the relationship between BEC and ICS effects is continuous; no / small effects are observed at lower BEC, with increasing effects at higher BEC.

The GOLD 2022 report has added additional evidence and recommendations concerning the 35 use of BEC in COPD in clinical practice. Notably, associations have been demonstrated in 36 COPD patients between higher BEC and increased levels of type-2 inflammation in the lungs. 37 These differences in type-2 inflammation can explain the differential ICS response according 38 to BEC. Additionally, lower BEC are associated with greater presence of proteobacteria, 39 notably haemophilus, and increased bacterial infections and pneumonia risk. These 40 observations support management strategies that use BEC to help identify subgroups with 41 42 increased ICS response (higher BEC) or increased risk of bacterial infection (lower BEC). Recent studies in younger individuals without COPD have also shown that higher BEC are 43 associated with increased risk of FEV₁ decline and the development of COPD. 44

45 Here we discuss and summarise the GOLD 2022 recommendations concerning the use of BEC46 as a biomarker that can facilitate a personalised management approach in COPD.

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52 Introduction

The Global Initiative for Chronic Obstructive Lung Disease (GOLD) published its first report 53 for the diagnosis and management of chronic obstructive pulmonary disease (COPD) in 2001 54 (1). Since then, GOLD has updated it yearly (2), the last time in 2022 (www.goldcopd.org). To 55 do so, GOLD evaluates critically the new evidence since the previous publication and decides 56 57 whether it merits (or not) inclusion in the most recent update. GOLD publishes specific recommendations and, sometimes, the main arguments behind them, but it often lacks space 58 for a detailed discussion regarding the pros and cons behind each recommendation. To address 59 60 this limitation, the Scientific Committee of GOLD decided to publish, separately from the main annual update, a series of papers that review and discuss topics of particular, current interest 61 62 for clinical practice.

The GOLD 2019 report recommended using blood eosinophil counts (BEC) as part of a precision medicine strategy to identify the most suitable patients for inhaled corticosteroid (ICS) treatment(3). Recent publications have provided further evidence and insights concerning BEC in COPD. Here, we discuss the role of BEC as a COPD biomarker, focusing on new advances and summarising the associated changes in the GOLD 2022 Report (shown in Table 1).

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70 A brief overview of eosinophil biology

Eosinophils originate from bone marrow stem cells, in response to stimulation by granulocytemonocyte colony stimulating factor (GM-CSF), interleukin (IL)-3 and IL-5 (4). The subsequent proliferation, activation, tissue infiltration and survival of eosinophils is controlled by type 2 (T2) mediators, such as IL-4, IL-5, IL-13 and eotaxins. Eosinophil degranulation releases major basic proteins, eosinophil cationic protein, eosinophil peroxidase and

eosinophil-derived neurotoxin, which provide host defence against parasitic infection (5). 76 These proteins also promote bacterial and viral clearance, although the extent of these roles in 77 humans, as opposed to animal models, is unclear (4, 5). Eosinophil derived granule proteins 78 can cause tissue injury and remodelling, while eosinophil peroxidase drives changes in the 79 physicochemical properties of mucus that underlie airway mucus plugging (4, 6). There is also 80 evidence that eosinophil subsets exist, with tissue resident cells having a predominantly 81 82 homeostatic role while inflammatory eosinophils are recruited into the lungs(7). Asthma and systemic hyper-eosinophilic diseases are examples where increased systemic and lung 83 84 eosinophil numbers, coupled with activation, contribute to disease pathophysiology (4).

85

86 BEC as a predictor of inhaled corticosteroid benefit

COPD is a heterogeneous condition, exemplified by the between individual variation in the 87 nature and severity of airway inflammation (3, 8-10). The use of anti-inflammatory treatments 88 therefore requires a selective approach, based on clinical characteristics (phenotyping) and 89 90 biological information (endotyping) in order to target therapies to subgroups of individuals who are most likely to derive benefit (3, 9, 11). ICS are anti-inflammatory drugs that are used 91 in combination with one or two long-acting bronchodilators (LABD) for the treatment of 92 93 COPD. Randomized control trials (RCTs) have shown that ICS reduce exacerbation rates, improve quality of life and prevent mortality when targeted to COPD patients with a history of 94 95 exacerbations (3, 12, 13). Pre-specified and post-hoc analysis of these RCTs have shown that higher BEC, used as a surrogate for lung eosinophil counts(14), at the study start are associated 96 with greater clinical benefits, notably exacerbation prevention, from ICS treatment (3, 14-17). 97 98 The relationship between BEC and ICS benefits has been described as continuous, as these analysis have demonstrated treatment effects at above (approximately) 100 cells / μ L with 99 incremental increases in the magnitude of effect at higher BEC(3, 14). Importantly, there is no 100

clear evidence that ICS treatment reduces BEC, so BEC retain their predictive value 101 independent of ICS treatment. Accordingly, in 2019 GOLD recommended the use of BEC in 102 clinical practice, in COPD patients with an exacerbation history despite appropriate use of 103 LABD, to identify the most suitable patients for ICS treatment (3). The BEC thresholds < 100 104 cells / μ L and \geq 300 cells / μ L have been proposed, identifying individuals with the lowest and 105 greatest likelihood (respectively) of benefit from ICS treatment when administered on top of 106 107 LABD. These are estimated, not strict, thresholds. Patients with low BEC appear to be at increased risk of pneumonia(18, 19) (discussed in depth later), while there is also a small 108 109 increase in pneumonia risk with ICS use in COPD patients(12, 13, 15).

RCTs of inhaled triple therapies have been analysed according to whether patients had 1 or ≥ 2 exacerbations in the previous year(20, 21). A history of ≥ 2 exacerbations was associated with more exacerbations during the study compared to 1 previous exacerbation. The benefit of ICS on exacerbation prevention was greater in individuals with more events (i.e. those with a history of ≥ 2 exacerbations), but there was still a benefit in patients with one previous exacerbation and BEC were able to predict ICS benefits regardless of exacerbation history.

Conclusion: The GOLD 2019 report recommended the use of clinical phenotyping (exacerbation history) combined with endotyping (using BEC as a biomarker) to enable ICS to be used with more precision, selecting individuals with a greater benefit (reduction in exacerbations) versus risk profile (pneumonia occurrence), hence increasing the net benefit potential of ICS (3). RCT results published since 2019 remain supportive of BEC as a predictive biomarker of ICS effects in COPD patients with increased exacerbation risk (15).

122 Variability of BEC

123 The intra-class correlation coefficient (ICC) for repeated BEC measurements performed on 124 different days in COPD patients have ranged from 0.64 to 0.89, indicating good to excellent

reproducibility(14). It has been commented that similar ICC values have been reported for
cholesterol and glycated Hb which are routinely used biomarkers in clinical practice(3, 14).
BEC show diurnal variation in healthy subjects and patients with asthma and COPD, peaking
in the early morning, and thought to be related to circadian variation in cortisol secretion(22,
23). The median reduction in BEC at 12.00 compared to 08.00 in COPD patients was reported
to be 36%(23).

GOLD has suggested BEC thresholds to help direct ICS treatment(3). Movement across a threshold after repeated measurement is more likely for BEC that are closer to the threshold(24). This is one reason why GOLD states that these are not strict thresholds, and consequently small within or between day variations should not result in a change in clinical management. In support, it has been reported that the predictive ability of BEC, with regard to ICS benefits observed in a triple therapy RCT, were similar regardless of whether the BEC at screening or randomisation was used, or the average of both(25).

138

BEC in COPD patients versus controls

A study in individuals aged >40 years showed that on average eosinophil counts were higher 140 in COPD patients (n=209) than in controls (n=127) (26). Although there was considerable 141 overlap in the counts between the groups, some COPD patients had higher counts than the 142 controls. A recent cohort study has also shown that BEC are higher in COPD patients (n=326) 143 versus controls (n=399)(27). In contrast, other studies have not shown differences between 144 COPD patients and controls (28), as the CANCOLD study showed a similar distribution of 145 146 BEC between the non-COPD (n=573) and COPD participants (n=547)(29), while the CHAIN cohort also showed a similar BEC distribution between non-COPD (n=121) and COPD 147 (n=769) participants(30). 148

A large general population study in Austria (n=11,042) using multivariate logistic regression 150 showed that a higher BEC (>210 cells/µL; the 75th percentile) was more likely in current 151 smokers (odds ratio (OR) 1.72, 95% confidence interval (CI)1.52-1.96) and COPD (OR 1.56, 152 CI 1.20–2.03), but the range in patients with COPD was not specified (31). In Japanese patients 153 with COPD (n=848), the median (interquartile range) BEC was 170 cells/µL (100-280 154 cells/µL) with a similar distribution to that in non-Japanese patients with COPD (n=5397), but 155 the counts were not compared to healthy controls(32). Another large general population study, 156 conducted in Japan (approximately 10,000 participants), showed a similar BEC distribution in 157 a healthy population to that seen in the European study, but BEC in patients with COPD were 158 not reported (33). 159

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A meta-analysis reported that the median BEC was higher in COPD patients compared to controls, although the 95% CI overlapped (34). There was high heterogeneity between studies, likely due to different characteristics of populations, particularly controls where co-morbid conditions that increase BEC (e.g. current smoking, allergies and obesity(31)) may not have been fully accounted for.

166

167 *Conclusion:* While the evidence is not consistent across all publications, there are three studies,
168 including a very large population study, showing that, on average, BEC are higher in COPD
169 patients, with a subgroup of COPD patients showing higher counts than seen in controls (26,
170 27, 31). These observations suggest upregulation of mechanisms that increase eosinophil
171 production from the bone marrow (i.e., the action of GM-CSF, IL-3 and IL-5(4)) or eosinophil

survival in some COPD patients. The lack of consistency across studies may reflect sample
size and / or the influence of co-morbidities on BEC (31).

174

175 BEC; association with future risk or disease progression

176 FEV1 decline

In healthy individuals who did not have asthma in the Dunedin Multidisciplinary Health and 177 Development Study (n=971), higher BEC were associated with faster FEV₁ decline between 178 the ages of 21 years and 38 years (35). The relationship persisted after adjusting for smoking. 179 Another study retrospectively analysed private healthcare screening records (n>359,000) of 180 younger adults without a history of asthma or airflow obstruction (mean age 36 yrs; median 181 182 follow up 5.6 years) (36). The development of airflow obstruction was associated with higher BEC at baseline, which was also observed in the smoker subgroup. Additionally, there was an 183 association between higher BEC and the development of physician diagnosed COPD plus 184 spirometric confirmation of airflow obstruction, defined as FEV1/FVC <0.7 and FEV1 <80%. 185 A limitation of this study is that post-bronchodilator spirometry was not performed. In the 186 187 CANCOLD study (n=1120; mean age 65 years), using a multivariate regression model which accounted for baseline factors including FEV₁, exacerbation history and ICS use, individuals 188 with BEC \geq 300 cells / μ L had more rapid FEV₁ decline than those with <150 cells / μ L (mean 189 190 difference 34.3 ml / year) (29). The same pattern was apparent in the COPD subgroup (n=466). Overall, these data from large cohort studies show that higher BEC are associated with more 191 rapid FEV₁ decline both in younger adults without airflow obstruction and patients with COPD 192 193 and, in some individuals, this leads to the development of COPD.

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Data from UK electronic medical records from COPD patients with $FEV_1 50 - 90\%$ predicted 195 (n=12,178) showed greater FEV₁ decline in patients with more exacerbations over >3 years 196 197 follow up (37). There was an interaction between exacerbation frequency and BEC, with a more rapid loss of lung function in patients with ≥ 2 exacerbations / year and BEC ≥ 350 198 cells/µL, which was reduced by ICS use. However, in patients without exacerbations, the rate 199 of FEV₁ decline was approximately 10ml/year less in patients with BEC≥350 cells/µL 200 201 compared to those with lower BEC. This study confirms the importance of exacerbations as a determinant of FEV₁ decline (38), and demonstrates complex relationships between BEC and 202 203 FEV₁ decline which are dependent on both exacerbation frequency and ICS use. An analysis of >26,000 COPD patients from the same database source showed that new ICS use versus no 204 ICS use was associated with reduced FEV1 decline in subjects with BEC >150 /uL(39), but 205 exacerbations were not analysed. A post-hoc analysis of the ISOLDE study also showed that 206 in patients with BEC $\geq 2\%$, FEV1 decline was reduced by ICS treatment (40). The Hokkaido 207 COPD cohort, with a smaller sample size (n=279) and low ICS use (<15%), reported that mean 208 BEC were lower in the rapid decliners compared to the slow decliners or sustainers (41). Again, 209 exacerbations were not reported in this study. 210

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Conclusion: In younger individuals without COPD, there is evidence of an association between 212 higher BEC and both faster FEV_1 decline and the development of airflow obstruction (35, 36). 213 These observations mechanistically implicate eosinophils and / or other associated components 214 of T2 inflammation in the development of COPD, at least in some patients. In patients with 215 216 confirmed COPD, the association between higher BEC and FEV₁ decline is complex and findings from cohort studies have been inconsistent, being influenced by disease heterogeneity 217 including prior exacerbation frequency and use of ICS(37-39). These complexities mean that 218 219 using BEC alone in COPD patients to predict lung function decline is a simplistic approach

that is unlikely to be of clinical utility. Nevertheless, FEV₁ decline appears to be greater in individuals with more exacerbations(37, 38), and ICS may reduce the rate of decline in individuals who have greater exacerbation risk plus higher BEC(37). These observational data, following COPD patients for \geq 3 years, support the results of multiple RCTs conducted over 1 year; both demonstrate a relationship between BEC and ICS benefits in COPD patients with a history of exacerbations (14-16, 42).

226

227 <u>Exacerbation risk</u>

Some cohort studies have reported an association between BEC and exacerbation risk in 228 patients with COPD, while others have found no relationship (30, 43-49). These contradictory 229 findings generally reflect differences in baseline exacerbation history (which is the strongest 230 predictor of exacerbation risk(50)) and ICS use, which RCTs have shown weaken the 231 relationship between exacerbation risk and BEC(14, 16, 17, 42). Cohort studies have generally 232 not adjusted for these factors. Analysis of two cohorts with prospective follow up data (n=1113 233 and n=1895) reported that BEC \geq 300 cells/µL were associated with increased exacerbation 234 frequency; this association was driven by the subgroup of individuals with ≥ 2 exacerbations in 235 the year before study start, with incident risk ratios of 1.96 and 1.4 for individuals with BEC \geq 236 $300 \text{ cells/}\mu\text{L}$ versus < $300 \text{ cells/}\mu\text{L}$ in this subgroup (51). The relationships between BEC and 237 exacerbation risk remained after adjusting for ICS use. 238

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Analyses of RCTs investigating ICS containing combination treatments in COPD patients with a history of exacerbations have shown that higher baseline BEC are associated with a higher rate of exacerbations over 12-months in patients not treated with ICS (15-17, 42). In contrast, a pooled analysis of 11 RCTs investigating LABD, involving patents with and without a history of exacerbations, found no relationship between BEC and exacerbation rates in patients not taking ICS (who also had lower exacerbation rates) (52). Exacerbation rates in patients taking ICS with were slightly higher (9%) in patients with BEC >300 cells / μ L compared to those with counts \leq 150 cells / μ L (52).

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Conclusion; Exacerbation history remains the best predictor of future exacerbation risk (50, 249 51). The potential usefulness of BEC as a predictor of future exacerbation risk is restricted to 250 patients with a history of exacerbations, and BEC have been consistently associated with 251 exacerbation risk in the non-ICS treatment arms of RCTs involving this clinical phenotype(15-252 17, 42). However, in cohort studies this relationship is less consistent, being modified by ICS 253 use and influenced by the inclusion of low exacerbation risk individuals(30, 43-49, 51). 254 Consequently, BEC are not a useful stand-alone biomarker of exacerbation risk in clinical 255 practice. 256

257

258 *Mortality*

In the CHAIN and BODE cohorts, all-cause mortality over 20 years was lower in COPD 259 260 patients with high BEC compared with those with values <300 cells/µL (15.8% versus 33.7%; p=0.026) after adjusting for age, sex, body mass index (BMI), lung function and Charlson 261 index(30). Over half the patients were taking ICS but the analysis was not adjusted for this. In 262 a French cohort, there was no relationship between BEC and 3-year survival, with over 85% 263 of patients taking ICS (46). The ETHOS RCT, conducted in patients at high exacerbation risk, 264 showed that the benefit of ICS (as part of triple combination treatment) on mortality was greater 265 at higher BEC(53). This mortality benefit due to ICS was accompanied by exacerbation 266 prevention at higher BEC(15). 267

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Conclusions. BEC used alone are not a reliable predictor of mortality, as the risk is modified
by ICS use. However, in the high exacerbation risk phenotype, RCT evidence supports higher
BEC as a biomarker of increased mortality risk in individuals not using ICS(53).

272

273 BEC and T2 inflammation

The consistent relationship between BEC and ICS effects on exacerbation rates in COPD RCTs 274 275 indicates that BEC reflect differential profiles of pulmonary inflammation within a heterogeneous condition(14, 16, 17, 54). Significant associations have been reported between 276 BEC and pulmonary eosinophil counts (from sputum or lung tissue), with the strength of the 277 278 relationship ranging from 0.18 to 0.7(49, 55-60). While these studies confirm that BEC reflect pulmonary eosinophil numbers, the association has been weak in some studies. The reasons 279 for a weak association include the inherent variability of lung sampling (e.g. between day 280 variation in sputum eosinophil counts(10)) and sometimes a lack of methodological precision 281 in eosinophil counts (e.g. using only one significant figure for BEC) (49). Furthermore, the 282 distribution of eosinophils in lung tissue is patchy (61), which may explain the lack of 283 association between blood and tissue eosinophils in one study (62), in contrast to the positive 284 relationship reported in other studies (55, 63, 64). 285

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Studies using bronchoscopy, induced sputum and lung surgical tissue samples have demonstrated a T2 inflammation profile in patients with higher BEC. Kolsum et al obtained bronchoscopy and sputum samples from 41 COPD patients with higher (>250 cells / μ L) or lower (<150 cells / μ L) BEC(64), and no previous asthma diagnosis or skin testing evidence of atopy. The higher BEC group had increased eosinophil counts in sputum, bronchoalveolar

lavage and bronchial mucosal tissue, plus increased protein levels of mediators involved in 292 eosinophil activation and chemotaxis (IL-5 and C-C motif chemokine ligand (CCL)24). The 293 higher BEC group also exhibited increased reticular basement membrane thickening. A 294 subsequent analysis of this study focused on gene expression of six T2 markers increased in 295 patients with asthma (65). Four genes, namely chloride channel accessory 1 (CLCA1), CCL26, 296 IL-13 and cystatin SN (CST1), had increased expression in both sputum cells and bronchial 297 298 brushings in the higher BEC COPD group, with these results validated in sputum samples from a different cohort (n=33). Bronchial epithelial brushings from the Emphysema versus Airway 299 300 disease (EvA) study (n=283) also showed differential gene expression in bronchial brushings from COPD patients with higher BEC, including CLCA1, CCL26 and CST1 (66). An asthma 301 cohort, analysed by the authors for comparison, showed far more differentially expressed genes 302 303 associated with BEC, suggesting a restricted T2 signature in COPD compared to asthma. Sputum cells obtained at the baseline visit of a RCT showed a differential gene expression 304 profile in samples with eosinophil counts $\geq 3\%$ versus < 3%, including known T2 markers(67). 305 Jogdand et al reported that eosinophil numbers in the conducting airways and lung parenchyma 306 were associated with more severe COPD and tissue basophil counts(61). 307

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Conclusion: Higher BEC in COPD patients are associated with both increased numbers of 309 eosinophils and levels of markers of T2 inflammation in the lungs(64-66). This differential 310 inflammation profile could explain the association between BEC and ICS responses, as T2 311 inflammation can respond well to corticosteroid treatment(68, 69). RCTs of biological 312 treatments targeting IL-5 or the IL-5 receptor, thereby reducing BEC, have failed to 313 demonstrate efficacy on exacerbation rates (the primary endpoint) in COPD populations 314 enriched for increased exacerbation risk and higher BEC(70, 71). A contributor to these 315 negative outcomes is that higher BEC appear to mark a wider T2 inflammation profile(61, 64-316

66), and selective depletion of eosinophil numbers will not modulate other T2 components.
BEC could be used as a biomarker to identify COPD patients suitable for clinical trials of novel
therapeutics targeting T2 pathways(14).

320

321 BEC and microbiome

Sputum samples obtained during the stable state from 510 patients with COPD were analysed 322 for cell counts and microbiome characteristics (by 16S rRNA sequencing) (10). Cross sectional 323 analysis showed that neutrophilic inflammation was associated with heterogeneous 324 microbiome patterns, including a subset with a haemophilus dominant microbiome. In contrast, 325 eosinophilic inflammation was associated with several non-dominant genera but not 326 haemophilus. Longitudinal analysis showed that eosinophilic samples that became non-327 eosinophilic over time also did not display a haemophilus dominant microbiome. Similarly, 328 studies in COPD patients using quantitative polymerase chain reaction quantification of 329 bacterial species have shown that haemophilus influenzae presence is associated with higher 330 sputum neutrophil counts and lower sputum eosinophil counts (72-74). Interestingly, 331 bronchoscopy samples from COPD patients with lower (versus higher) BEC showed decreased 332 immunoglobulin subtype levels and reduced opsonisation of non-typeable haemophilus 333 influenzae; this provides a possible explanation for higher sputum haemophilus influenzae 334 levels in patients with lower eosinophil counts (75). 335

336

337 Dicker et al showed that higher BEC were associated with lower proteobacteria abundance 338 (which includes the haemophilus genera), and greater abundance of the firmicutes phyla in a 339 cohort of 296 COPD patients(76). Furthermore, there was an increase in haemophilus 340 abundance for patients with BEC ≤ 100 cells / μ L compared to > 100 cells / μ L. Subgroup analysis showed that the profile of inflammatory proteins in sputum was different in samples
with proteobacteria dominance, favouring mediators of neutrophilic inflammation, when
compared to firmicutes dominant samples. Overall, these cohort studies have highlighted that
lower eosinophil counts (in sputum and blood) are associated with a different microbiome
profile, characterised by increased proteobacteria.

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Martinez-Garcia et al reported that BEC<100 cells / µL were associated with increased 347 incidence of chronic bacterial infection (CBI) and pneumonia episodes in 201 COPD patients 348 (median follow up 7 years) (19). A multivariate regression model showed that age, FEV_1 , CBI, 349 and BEC<100 cells/µl were all independently associated with greater pneumonia risk. Higher 350 BEC thresholds (<150 cells / μ l and <300 cells / μ l) were not significantly associated with 351 increased pneumonia risk. ICS use was not associated with pneumonia risk in the overall 352 population, although ICS further increased the risk of pneumonia (hazard ratio [HR] 2.9) in 353 those with CBI and <100 eosinophils/µl. A pooled analysis of 10 randomised control trials of 354 ICS containing combination treatments in COPD patients showed that the risk of pneumonia 355 was higher in patients at baseline BEC <2% versus \geq 2%; HR 1·31 (95% CI 1·06–1·62)(18). A 356 potential explanation for these pneumonia findings comes from a small COPD RCT (n=60) 357 which showed that ICS containing combination treatment over one year increased sputum 358 bacterial load, in contrast to no change without ICS; this increase was present in those with 359 lower BEC only(77). 360

361

362 *Conclusion:* Recent studies have consistently shown that lower sputum and blood eosinophil 363 counts are associated with an increased presence of proteobacteria phylum/haemophilus 364 genera(10, 72-74, 76). Lower BEC also appear to be associated with an increased risk of recurrent bacterial infections and pneumonia, and these risks seem to be increased by ICS use in patients with lower BEC(19, 77, 78). Overall, these findings regarding microbiome and pneumonia risk provide additional reasons not to use ICS in COPD patients with lower BEC.

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369 Summary and conclusions

The GOLD 2019 report first introduced BEC as a biomarker to help make pharmacological 370 treatment decisions, concerning ICS use, in COPD patients with a history of exacerbations (3). 371 The GOLD 2022 report now adds various additional evidence concerning BEC (key points 372 shown in Table 1), including the connections between BEC, T2 inflammation(61, 64-66) and 373 lung microbiome(10, 72-74, 76) which identify COPD subgroups with increased ICS response 374 (higher BEC) or increased risk of bacterial infection (lower BEC); summarised in Figure 1. 375 This evidence supports an integrated evaluation of clinical history (notably exacerbation 376 history), BEC and sputum microbiology in order to provide a personalised management 377 approach with regard to when ICS should be used on top of LABD and the management of 378 airway infection. 379

Accumulating evidence indicates an association between lower BEC and the incidence of both 380 CBI and pneumonia events (18, 19), coupled with a differential microbiome profile (greater 381 abundance of haemophilus influenza)(10, 72-74, 76). Based on this evidence, lower BEC (<100 382 cells/µl) could be used as a biomarker, in combination with clinical history, to help identify 383 patients who require careful monitoring for bacterial colonisation. Furthermore, in these 384 individuals, the absence of T2 inflammation coupled with the increased risk of bacterial 385 386 infection, argues against the use of ICS. The importance of bacterial colonization was demonstrated in an observational COPD cohort where exacerbation risk was greatest in 387 individuals with haemophilus influenza colonization and exposure to rhinovirus infection (79), 388

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indicating an interplay between pathogens leading to worse clinical outcomes. Further studies
should elucidate the mechanisms responsible for the association between T2 inflammation and
the microbiome, as this may help identify novel therapeutic interventions.

392

COPD patients with higher BEC have more T2 inflammation (61, 64-66), which can explain a 393 differential response to ICS. It is important to note that RCTs have demonstrated a benefit for 394 ICS (as part of combination treatments) only in COPD patients with an exacerbation history in 395 the previous year (3, 14). There is currently no evidence supporting ICS intervention in COPD 396 patients with higher BEC but without a history of exacerbations, although this is an evidence 397 gap worth considering. Furthermore, the association between higher BEC and FEV1 decline in 398 younger adults (36) provides a rationale to study the effects of ICS on disease progression / 399 lung function decline in younger COPD patients with higher BEC (80). 400

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BEC are not a stand-alone biomarker of future risk (of FEV1 decline, exacerbations, and mortality) in patients with COPD, due to the complex relationship with exacerbation risk and confounding due to ICS use(37). However, in younger individuals, higher BEC may serve as a biomarker to help identify those at increased risk of developing COPD (36), and further evidence is needed to evaluate the utility of BEC in this context.

RCTs have shown that, in COPD patients with a history of exacerbations, higher BEC identify a subgroup with increased exacerbation risk that can be therapeutically modified by ICS(15-17). On the other hand, we also point out a subgroup with lower BEC (<100 cells/µl) with a different microbiome profile and increased risk of chronic bacterial infection(19, 76). These findings might suggest that BEC predict a "U shaped" future risk curve, albeit one that is influenced by other factors including exacerbation history and ICS use. In conclusion, the GOLD 2022 report incorporates new evidence regarding BEC, notably the

relationships with T2 inflammation(64-66) and the microbiome(10, 72-74, 76). These findings

415 further our understanding of COPD subtypes, facilitating precision medicine strategies based

416 on clinical phenotyping combined with endotyping(9, 11).

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422

423 Figure legends

424 Figure 1: The relationships between blood eosinophil counts (BEC) and Type-2 (T2)

inflammation, microbiome, bacterial infection / pneumonia episodes and inhaled corticosteroid

426 (ICS) response (exacerbation prevention).

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to Review Only

Table 1. GOLD 2022 Report: Key evidence and recommendations

for blood eosinophil counts in COPD

Prediction of ICS benefits

- The use of BEC to predict ICS effects should be combined with exacerbation risk (using exacerbation history)
- The relationship between BEC and ICS effects is continuous; no / small effects are observed at lower BEC, with increasing effects at higher BEC
- < 100 cells/ μ L and \geq 300 cells/ μ L are estimates, not precise cut-off values, to identify individuals with the lowest and greatest (respectively) likelihood of ICS benefit

Type-2 inflammation

- Higher BEC are associated with increased lung eosinophil numbers and higher levels of type-2 inflammation markers in the airways
- The differences in type-2 inflammation can explain the differential ICS response according to BEC

COPD versus controls

• A subset of COPD patients have BEC above those found in controls

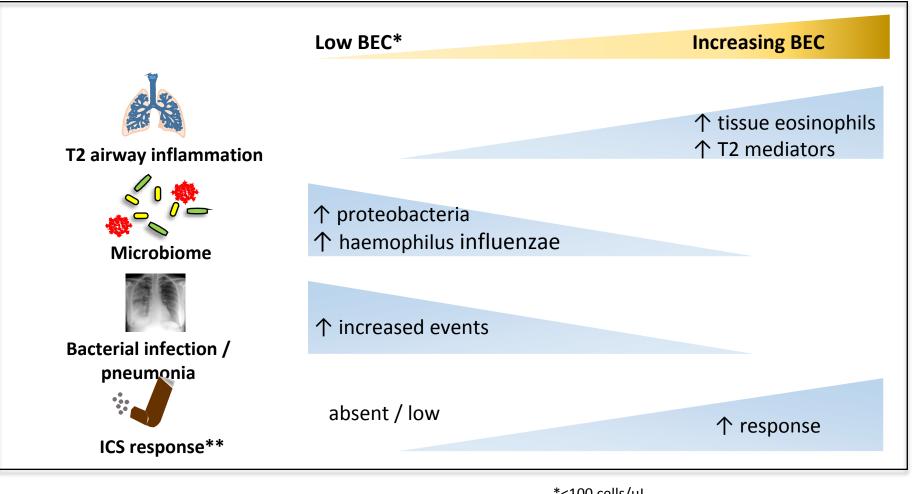
Microbiome

• Lower BEC are associated with greater presence of proteobacteria, notably haemophilus, and increased bacterial infections and pneumonia

Future risk (of exacerbations / disease progression)

- In younger individuals without COPD, higher BEC are associated with increased risk of FEV₁ decline and the development of COPD
- BEC cannot be used as a standalone biomarker of future risk without considering exacerbation risk and ICS use

Abbreviations: BEC = blood eosinophil count. ICS= inhaled corticosteroid



*<100 cells/μL **In COPD patients with increased exacerbation risk