

VIEWPOINT

# Could lobar flow sequencing account for convection-dependent ventilation heterogeneity in normal humans?

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INDICES OF CONVECTION-DEPENDENT ventilation heterogeneity have been used as a means to noninvasively sample airway structure down to the entrance of the gas exchanging zone. Although potentially attractive to reflect abnormal function of small nonalveolated airways or entire tissue units as small as acini, these indices can also be affected by much larger-scale ventilation heterogeneity. For instance in normal humans, gravity causes greater specific ventilation in the dependent parts of the lungs in the line of gravity (6, 12). Irrespective of the scale at which it is generated, convection-dependent concentration heterogeneity produces a curvilinear washout at the mouth, i.e., a curvilinear semilog plot of breath-by-breath mean expired concentration of a multiple breath washout (MBW)

(22). This can be quantified by a nonzero Curv (23), with a maximum value of 1, which corresponds to the presence of an infinitesimally slow lung unit. If in addition to a nonzero Curv, the best ventilated lung units are the ones to contribute relatively more to expiratory flow in the early vs. late phase of each expiration, this produces a positive N<sub>2</sub> phase III slope that shows a breath-by-breath increase when normalized by expired concentration (2); this increase is most often characterized by the index Scnd (18, 23). It follows that a nonzero Scnd always implies that Curv is nonzero too, although a nonzero Curv can be associated with zero Scnd (in the absence of flow sequencing, i.e., all flow proportions constant during expiration). MBW experiments in microgravity and in different body postures (16, 17, 19) can lead to different Curv and Scnd values, but in normal humans both indices are nonzero and Scnd (and the underlying N<sub>2</sub> phase III slope) is always positive.

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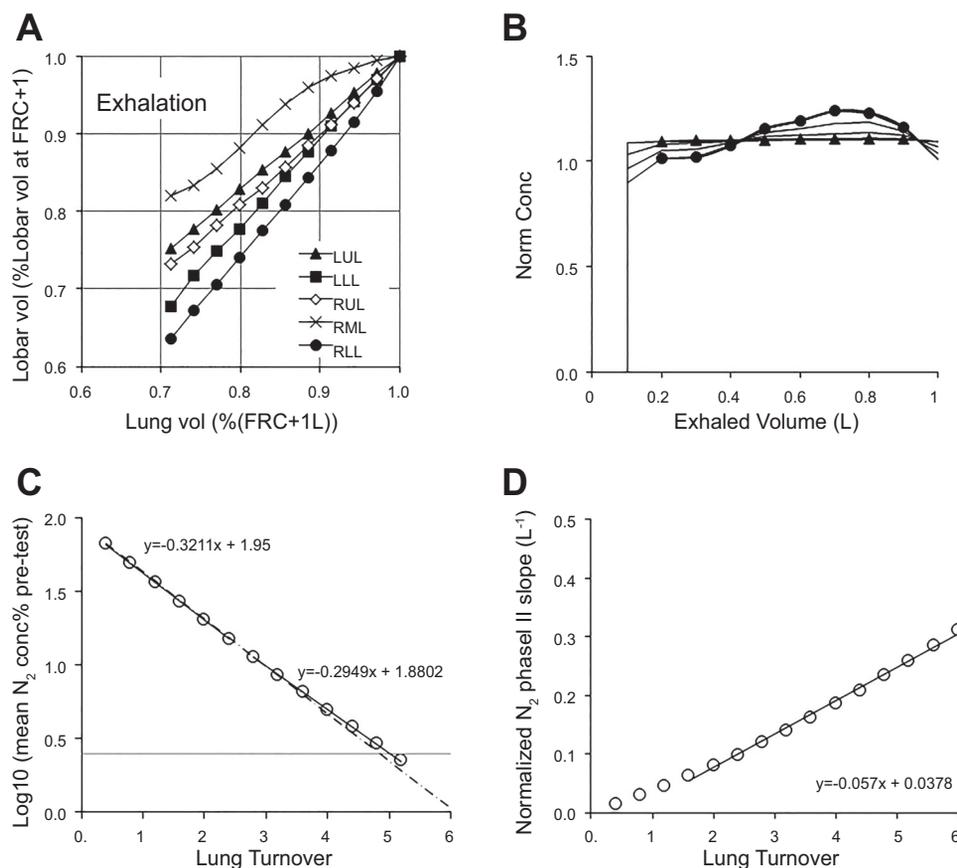


Fig. 1. A: expiratory lobar volume vs. total lung volume, as a percentage of respective end-inspiratory volumes, for the 5 lobes according to the digitization of Fig. 5 in Jahani et al. (8) (considering respiration between FRC and FRC+1l). B: N<sub>2</sub> concentration traces for breath number 1, 5, 10, and 15 (normalized to mean expired N<sub>2</sub> concentration over the entire breath including dead space), resulting from each subsequent O<sub>2</sub> inspiration according to the lobar volume distribution in A (considering 5 lobar compartments, FRC = 2.5 liters and 100 ml dead space). ▲ (breath 1) and ● (breath 15) show the concentrations that were used for calculation of phase III slope regression within each expiration. C and D: washout simulations of N<sub>2</sub> gas concentration and normalized N<sub>2</sub> phase III slopes vs. lung turnover, resulting from each subsequent O<sub>2</sub> inspiration according to the lobar volume distribution in A (considering 5 lobar compartments, FRC = 2.5 liters and 100 ml dead space); uniform mixing is also represented (dashed dotted line). The corresponding simulated Curv = 0.08 (= 1-0.2949/0.3211; one minus the regression slope ratio over the 2 halves of the turnover range down to 1/40th initial concentration in the washout concentration curve indicated by the grey horizontal line) and simulated Scnd = 0.057 liter<sup>-1</sup> (regression slope of normalized N<sub>2</sub> phase III slope between 1.5 and 6 lung turnovers).

Based on data obtained from radiopaque parenchymal markers in dog lungs (7, 14), Wilson et al. (27) suggested that the N<sub>2</sub> phase III slope observed in excised dog lobes could be due to increased ventilation variability at a small (intraregional) scale. At least two intraregional mechanisms, parenchymal expansion dynamics (26) and airway-parenchyma interdependence (10), have the potential to generate both conditions for a nonzero phase III slope: specific ventilation heterogeneity and expiratory flow sequencing. Thus far it has been impossible to demonstrate that either mechanism alone or embedded in a realistic airway network (13) would automatically, and without any parameter fitting, lead to a positive simulated N<sub>2</sub> phase III slope in normal humans. When Young and Martin investigated the role of lobar dynamics on ventilation heterogeneity by means of challenging bronchoscopic experiments (9, 11, 28), they reported flow or concentration measurements between lobes in a single lung to specifically assess upper and lower lung zones (9) or to infer that interlobar effects on concentration traces are overridden by intralobar effects (28). In the meantime, these intralobar effects can be accounted for in large part by diffusion-convection-dependent ventilation heterogeneity (15). Modeling and experimental studies (15, 25) have shown why diffusion-convection interaction in the lung periphery always generates a positive N<sub>2</sub> phase III slope. As for the intraregional convection-dependent part of the N<sub>2</sub> phase III slope in normal humans that was already the subject of a literature study 30 years ago (4), the jury is still out. Based on recent lung imaging data, the present Viewpoint explores the possibility that the interplay between all five lung lobes could in fact produce a positive N<sub>2</sub> phase III slope and therefore at least partly account for Scond in normal humans.

A dynamic CT study on 6 healthy subjects aged 24–58 years (8) measured lobar volumes at 20 intervals within tidal breathing cycles. In Fig. 1, we plotted these lobar volumes vs. total lung volume, resembling the familiar onion skin diagram (12), but adapted to a 1-liter volume excursion starting from FRC (and showing only the expiratory volume excursions) (Fig. 1A). This shows that with the subjects supine in the CT, the diaphragmatic lobes are still the ones being least expanded at FRC and best ventilated. This alone will generate a nonzero Curv. In addition, the curvilinear plots in Fig. 1A signal flow sequencing that can also generate a nonzero Scond. Both Curv and Scond can be simulated as follows. Using Fig. 1A, end-inspiratory concentrations in each lobe can be computed and recombined according to expiratory flows from each lobe to simulate a washout curve (23) (Fig. 1, B–D); this obtains Curv = 0.08 and Scond = 0.057 liter<sup>-1</sup>. By comparing these simulated values to average experimental data for a 40-yr-old normal subject (Curv = 0.18 and Scond = 0.032 liter<sup>-1</sup>) (24), we observe that specific ventilation heterogeneity (affecting Curv and Scond) is too small and flow sequencing (affecting only Scond) is too big to match normal conductive ventilation heterogeneity. Calculations would need to be repeated on per-subject sets of dynamic lobar flow curves to also assess intersubject variability. Nevertheless, the importance of this small lobar data set (8) is that it reveals a pattern of lobar flow sequencing that is compatible with a nonzero, and indeed positive, N<sub>2</sub> phase III slope.

In a larger group of almost 100 normal never-smoker subjects aged 45–80 yr lobar volumes at FRC and near TLC were obtained by static CT (29). If in a first approximation, we

assume that relative lobar volume changes between FRC and FRC+I1 are proportional to those between FRC and TLC, this results in simulated Curv = 0.19 (male) and 0.25 (female) [average experimental Curv = 0.24 for a 60-yr-old normal subject (24)]. Depending on the degree of nonlinearity in lobar volume excursions between FRC and TLC, the estimate of Curv will be affected. However, dynamic CT studies measuring such nonlinearity (8) cannot be readily reproduced in a large healthy population for ethical reasons. Ventilation modalities that visualize inhaled tracer concentration by scintigraphy (5) or MRI (1) would also require a delineation of lobes, for instance by registration with CT images at some instances in the breathing cycle. Still, this might not obtain the appropriate accuracy to distinguish a real difference in relative lobar specific ventilation from its measurement noise; the latter would also lead to a simulated nonzero Curv. An attractive alternative is the measurement of velocity profiles over the cross sections of the main bronchi by MRI phase-contrast velocimetry (1), which has been able to show that at very low flows and in the supine position, there is a cardiogenic pendelluft between the left and right lung lobes (1, 21). Despite considerable intersubject variability, the lobar expansion and deflation was seen to be most frequently influenced by local volume changes in synchrony with the heart beat in the left lower lobe (1). Potentially, this technique could be elaborated further to also obtain velocity over the five bronchi leading to the five lobes and derive lobar flows during a full breathing cycle.

We conclude this Viewpoint by contending that lobar expansion is a likely contributor to the convection-dependent N<sub>2</sub> phase III slope and that an adaptation of noninvasive imaging (1) or reanalysis of existing CT data (8) could help confirm or refute this hypothesis. Because the recumbent body posture during imaging likely affects functional residual capacity (3) and lobar expansion, these imaging data strictly apply only to MBW tests performed in the same posture. A bronchoscopic system currently used to detect collateral ventilation in chronic obstructive lung patients eligible for lobar resection (20) could be modified to obtain relative lobar flows in upright humans, provided flow cycles measured in the five different lobar bronchi can be properly aligned to quantify flow sequencing between them.

#### DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the author(s).

#### AUTHOR CONTRIBUTIONS

S.A.V. drafted manuscript; S.A.V. and M.P. edited and revised manuscript; S.A.V. and M.P. approved final version of manuscript.

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