

S1 File: Supplement

Effect of dexamethasone exposure on the neonatal unit on adolescent lung function

Christopher Harris, Siobhan Crichton, Sanja Zivanovic, Alan Lunt, Sandy Calvert, Neil Marlow, Janet L Peacock, Anne Greenough

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Further Baseline data (S1 file: Online supplement. Tables A-C)

Table A shows lung function at follow up in children with or without complete data and shows little evidence of bias in our dataset.

Table B shows reasonable balance according to whether follow-up lung function data were or were not available.

Table C gives mean FEF₇₅ z-score by neonatal factors and shows that most neonatal factors had a limited effect on later lung function. This supports our suggestion that the observed effects of dexamethasone are unlikely to be fully explained by uncontrolled confounding.

26 S1 File: Online supplement. Table A: Lung function at follow up of children with and without
 27 complete data.

		Complete covariate data	Missing covariate data	
Lung Function	N	Mean(SD) (n=179)	Mean(SD) (n=69)	p-value
FEF ₇₅ z score	248	-1.09 (0.89)	-1.03 (0.87)	0.748
FEF ₅₀ z score	248	-1.22 (0.92)	-1.20 (0.85)	0.837
FEF ₂₅ z score	248	-1.02 (0.95)	-0.95 (0.89)	0.403
FEF ₂₅₋₇₅ z score	231	-1.46 (1.11)	-1.46 (0.97)	0.995
FEV ₁ z score	248	-0.80 (1.10)	-0.71 (0.95)	0.563
FVC z score	248	-0.38 (1.02)	-0.32 (0.84)	0.769
FEV ₁ :FVC z score	248	-1.49 (1.88)	-1.33 (1.50)	0.527
PEF percentage predicted	247	83.6 (14.9)	82.9 (16.3)	0.991
RV z score	211	0.54 (1.35)	0.06 (0.96)	0.018
FRC _{pleth} z score	218	0.03 (1.31)	-0.40 (1.11)	0.034
FRC _{he} z score	229	-0.64 (1.07)	-0.77 (1.06)	0.469
DL _{CO} z score	210	-0.94 (1.09)	-0.97 (1.03)	0.941
Respiratory resistance - percentage predicted				
At 5 Hz	237	97.2 (23.0)	91.3 (19.6)	0.098
At 20 Hz	237	93.2 (23.0)	89.76 (22.5)	0.380

28 S1 file: Online supplement. Table B: Baseline characteristics of the infants included and not included due to missing lung function data
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	Sample analysed with complete data N=179	Sample not analysed with incomplete data N=223	Comparison of complete and incomplete samples
Characteristics	% (n) or mean (SD)	% (n) or mean (SD)	p-value
Birth weight	882 (208)	912 (211)	0.15
Gestational age	26.5 (1.3)	26.2 (1.4)	0.06
Male sex	51% (91)	54% (125)	0.57
Multiple birth	25% (44)	22% (52)	0.59
Postnatal dexamethasone use	28% (50)	30% (70)	0.64
Oxygen dependency at 28 days	82% (147)	81% (188)	0.71
Oxygen dependency at 36 weeks PMA	59% (105)	58% (135)	0.88
Oxygen dependency at hospital discharge	24% (43)	21% (50)	0.50
Major ultrasound abnormality in neonatal period	13% (23)	15% (36)	0.46
Maternal smoking in pregnancy	23% (41)	27% (58)	0.31

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31 S1 File: Online supplement. Table C: Mean FEF₇₅ z-score by neonatal factors (n=179)

	Mean FEF₇₅ z-score (SD)	p-value
Birth weight <860g (89) ≥860g (90)	-1.12 (0.95) -1.07 (0.83)	0.71
Birthweight standard deviation score <-0.5 (90) ≥-0.5 (89)	-1.16 (0.86) -1.02 (0.91)	0.28
Gestational age 23-25wk (42) 26-28wk (137)	-1.11 (0.95) -1.09 (0.87)	0.91
Sex Girl (88) Boy (91)	-1.00 (1.03) -1.18 (0.71)	0.18
Multiple birth Singleton (135) Multiple (44)	-1.22 (0.82) -0.70 (0.98)	<0.001
Oxygen dependency at 36 weeks PMA No (74) Yes (105)	-0.95 (0.84) -1.19 (0.91)	0.08
Neonatal cranial ultrasound Normal (156) Abnormal (23)	-1.09 (0.91) -1.11 (0.72)	0.92
Airleak No (158) Yes (21)	-1.10 (0.89) -1.02 (0.87)	0.68
Patent ductus arteriosus No (125) Yes (54)	-1.11 (0.92) -1.05 (0.80)	0.72
Pulmonary haemorrhage		

No (169)	-1.07 (0.89)	0.15
Yes (10)	-1.48 (0.85)	
Mode of ventilation		
CV (90)	-1.19 (0.79)	0.13
HFOV (89)	-0.99 (0.96)	
Apgar score at 5 mins		
<9 (82)	-1.06 (0.95)	0.66
≥9 (97)	-1.12 (0.83)	
Maternal smoking in pregnancy		
No (138)	-1.01 (0.86)	0.65
Yes (41)	-1.04 (0.99)	
Antenatal steroids		
No (17)	-1.09 (0.67)	0.98
Yes (161)	-1.09 (0.91)	

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34 **Propensity Score Matching (S1 File: Online supplement. Table D)**

35 Since the use of postnatal dexamethasone is so highly confounded by neonatal factors, we also used propensity score matching for
36 dexamethasone exposure (yes/no) [S4] as an alternative way of adjustment in addition to linear mixed model regression. Propensity score (PS)
37 matching works differently to multiple regression in that it matches the subjects as closely as possible using baseline factors prior to analysis so
38 that the study closely resembles a randomised trial

39 The matching algorithm used was up to three nearest neighbours for each case. This approach was used because it leads to less bias than if the
40 single nearest neighbour approach is used without replacement [S4]. Logistic regression models were used to assign a probability of steroid use
41 to each child based on their baseline characteristics. The model included data collected prior to the initiation of steroid use, namely: sex, birth
42 weight, birth weight z-score, gestational age in weeks (in keeping with the original trial's randomisation strata), smoking in pregnancy, multiple
43 birth, ventilation group and Apgar score at five minutes. Children who received dexamethasone, were then matched to three children who did not
44 receive dexamethasone with the closest propensity scores.

45 The main challenge of the PS method is to obtain close matches for all subjects. Inspection of the table of variables by groups before and after
46 matching showed substantial improvement achieved by PS with no significant imbalance for any variable. It was not possible to use propensity
47 score matching for three measures of dexamethasone exposure, that is timing of administration, number of courses and days of exposure due to
48 the small numbers in the different dexamethasone-use categories. For this reason, only adjustment by multivariable logistic regression was

49 undertaken for those measures. A further limitation of PS matching in this context is that it is difficult to adjust for clustering within propensity
50 score models.

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52 S1 File: Online supplement. Table D: Lung function and postnatal dexamethasone exposure: sensitivity analyses adjusted for confounding using propensity
 53 score matching
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		No dexamethasone exposure	Dexamethasone exposure	Adjusted using multiple regression (main analysis)		Adjusted using propensity score matching (sensitivity)	
Lung Function	N	Mean (SD)	Mean (SD)	Difference (95% CI)	p-value	Difference (95% CI)	p-value
FEF ₇₅ z score	179	-0.95 (0.91)	-1.45 (0.71)	-0.53* (-0.85 to -0.21)	0.002	-0.51 (-0.89 to -0.13)	0.009
FEF ₅₀ z score	179	-1.04 (0.89)	-1.71 (0.81)	-0.74 (-1.05 to -0.43)	<0.001	-0.54 (-0.93 to -0.14)	0.006
FEF ₂₅ z score	179	-0.82 (0.91)	-1.53 (0.86)	-0.75 (-1.07 to -0.44)	<0.001	-0.51 (-0.78 to -0.24)	<0.001
FEF ₂₅₋₇₅ z score	169	-1.24 (1.07)	-1.98 (1.05)	-0.70 (-1.08 to -0.33)	<0.001	-0.55 (-0.99 to -0.11)	0.014
FEV ₁ z score	179	-0.55 (1.03)	-1.44 (1.03)	-0.87 (-1.24 to -0.51)	<0.001	-0.62 (-1.00 to -0.24)	0.002
FVC z score	179	-0.24 (0.96)	-0.73 (1.11)	-0.38 (-0.75 to -0.01)	0.043	-0.23 (-0.59 to 0.14)	0.221
FEV ₁ :FVC z	179	-1.17 (1.69)	-2.32 (2.11)	-1.43 (-2.09 to -0.78)	<0.001	-1.16 (-1.98 to -0.34)	0.006
PEF % pred*	178	86.07 (14.64)	77.36 (13.98)	-10.74 (-16.06 to -5.41)	<0.001	-7.42 (-11.5 to -3.4)	<0.001
RV z score	152	0.26 (1.09)	1.29 (1.67)	0.86 (0.36 to 1.36)	0.001	0.67 (0.27 to 1.07)	0.001
FRC _{pleth} z	157	-0.11 (1.25)	0.39 (1.39)	0.39 (-0.11 to 0.90)	0.128	0.37 (0.01 to 0.73)	0.042
FRC _{he} z score	168	-0.73 (1.09)	-0.42 (1.00)	0.27 (-0.13 to 0.66)	0.186	0.30 (-0.06 to 0.67)	0.106
DL _{CO} z score	149	-0.93 (1.11)	-1.04 (1.02)	0.09 (-0.33 to 0.52)	0.658	-0.01 (-0.61 to 0.59)	0.968
At 5 Hz	170	96.06 (21.38)	100.11 (27.03)	9.57 (1.13 to 18.02)	0.026	1.22 (-5.42 to 7.86)	0.719
At 20 Hz	170	93.94	91.28	2.49	0.578	-1.57	0.669

		(19.82)	(4.46)	(-6.27 to 11.25)		(-8.79 to 5.64)	
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55 **Further details on statistical analysis (S1 File: Online supplement. Tables E**
56 **and F)**

57 Since differences in mean z-scores can be difficult to interpret, we have additionally
58 presented the equivalent difference in the proportion of children with abnormal lung function.
59 To do this we have used the 5th centile for normal to define the cut-off between ‘normal’ and
60 ‘abnormal’. Since in healthy children the z-score has a Normal (Gaussian) distribution, the
61 cut-point for abnormality is defined as $z < -1.645$, the 5th centile of the Normal Distribution.

62 The proportion abnormal is not calculated using the data values themselves but using a
63 statistical model to gain precision. These calculations are similar to those performed to
64 calculate reference ranges. The calculations used a statistical method called the
65 ‘distributional approach’ [table 3] The distributional approach provides more precise values
66 than we would obtain had we used the data alone. In the present study, the adjusted estimates
67 from the multivariable mixed model analyses were used to estimate the difference in the
68 proportion of children who have abnormal lung function in those children who were and
69 were not exposed to steroids. The calculations use the same adjusting factors to allow for
70 confounding neonatal factors as the main analyses [table 3].

71 The differences for all analyses in main paper Table 2 are shown in Table 3. They show that
72 what may seem to be quite small differences in mean z-score, eg, 0.53 standard deviations for
73 FEF₇₅, translate into quite substantial differences in the percentage that have abnormal lung
74 function results, 22 percentage points, and so the additional data help to make the results
75 more clinically meaningful.

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77 Table S5 is a further sensitivity analysis that adjusts for antenatal steroids and postnatal
78 surfactant. This table shows no appreciable effect of this change in the modelling. Table S6
79 gives the random effects estimates for the mixed effects models shown in the main paper,
80 table 2.

S1 File: Online supplement. Table E: Sensitivity analyses adjusting for antenatal steroids and postnatal surfactant

		No dexamethasone exposure	Dexamethasone exposure	Adjusted using original variables (Table 2 in text)	Adjusted using original variables + antenatal steroids, postnatal surfactant
Lung Function	N	Mean (SD)	Mean (SD)	Difference (exposed-unexpo) (95% CI)	Difference (exposed-unexpo) (95% CI)
FEF ₇₅ z score	179	-0.95 (0.91)	-1.45 (0.71)	-0.53 (-0.85 to -0.21)	-0.52 (-0.84 to -0.20)
FEF ₅₀ z score	179	-1.04 (0.89)	-1.71 (0.81)	-0.74 (-1.05 to -0.43)	-0.73 (-1.04 to -0.43)
FEF ₂₅ z score	179	-0.82 (0.91)	-1.53 (0.86)	-0.75 (-1.07 to -0.44)	-0.77 (-1.09 to -0.46)
FEF ₂₅₋₇₅ z score	169	-1.24 (1.07)	-1.98 (1.05)	-0.70 (-1.08 to -0.33)	-0.68 (-1.06 to -0.31)
FEV ₁ z score	179	-0.55 (1.03)	-1.44 (1.03)	-0.87 (-1.24 to -0.51)	-0.88 (-1.24 to -0.51)
FVC z score	179	-0.24 (0.96)	-0.73 (1.11)	-0.38 (-0.75 to -0.01)	-0.40 (-0.77 to -0.02)
FEV ₁ :FVC z	179	-1.17 (1.69)	-2.32 (2.11)	-1.43 (-2.09 to -0.78)	-1.42 (-2.08 to -0.75)
PEF % pred*	178	86.07 (14.64)	77.36 (13.98)	-10.74 (-16.06 to -5.41)	-10.75 (-16.15 to -5.35)
RV z score	152	0.26 (1.09)	1.29 (1.67)	0.86 (0.36 to 1.36)	0.86 (0.38 to 1.39)
FRC _{pleth} z	157	-0.11 (1.25)	0.39 (1.39)	0.39 (-0.11 to 0.90)	0.40 (-0.12 to 0.91)
FRC _{he} z score	168	-0.73 (1.09)	-0.42 (1.00)	0.27 (-0.13 to 0.66)	0.21 (-0.18 to 0.61)
DL _{CO} z score	149	-0.93 (1.11)	-1.04 (1.02)	0.09 (-0.33 to 0.52)	0.06 (-0.36 to 0.49)
At 5 Hz	170	96.06	100.11	9.57	8.60

		(21.38)	(27.03)	(1.13 to 18.02)	(0.10 to 17.10)
At 20 Hz	170	93.94 (19.82)	91.28 (4.46)	2.49 (-6.27 to 11.25)	1.84 (-7.02 to 10.71)

- 1 S1 File: Online supplement. Table F: Random effects estimates from adjusted models presented in
- 2 table 2 main paper

Lung Function	SD Intercept	SD Residual
FEF ₇₅ z score	0.54	0.62
FEF ₅₀ z score	0.60	0.54
FEF ₂₅ z score	0.52	0.63
FEF ₂₅₋₇₅ z score	0.80	0.55
FEV ₁ z score	0.58	0.74
FVC z score	0.58	0.76
FEV ₁ :FVC z score	1.25	1.19
PEF % pred*	10.96	8.55
RV z score	0.70	0.93
FRC _{pleth} Z	0.80	0.97
FRC _{he} z score	0.62	0.78
DL _{CO} z score	0.73	0.70
At 5 Hz	16.41	14.14
At 20 Hz	15.42	16.13

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5 **References**

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