

Attention deficit hyperactivity disorder (update)

**Appendix 3: Cost-effectiveness analysis:
Network meta-analysis for ADHD treatments in
combination and individually**

NICE guideline CG72

Economic analysis report

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*This guideline was developed by the
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1 **Cost-effectiveness analysis: Network meta-analysis for ADHD treatments in combination and individually**

4 **1.1 Introduction**

5 The results of conventional meta-analyses of direct evidence alone (as presented in the
6 GRADE profiles and forest plots in evidence review F on combination treatment) does not
7 help inform which intervention is most effective for managing the symptoms of ADHD. The
8 challenge of interpretation has arisen for two reasons:

- 9 • In isolation, each pair-wise comparison does not inform the choice among the different
10 treatments because there are more than two treatments being compared in the
11 combination review. In addition direct evidence is not available for some pair-wise
12 comparisons in a randomised controlled trial.
- 13 • There could be conflicting estimates of effect if we try to compare the results of different
14 pairwise comparisons if trying to decide which intervention is best.

15 An additional problem is that the clinical data needed in a model is dichotomous in nature
16 (because of needing to link to quality of life data), whereas the clinical review focused on
17 continuous outcomes primarily, and therefore to weigh up the costs, benefits and harms of
18 the different interventions additional analysis of the data is needed.

19 To overcome these problems, a hierarchical Bayesian network meta-analysis (NMA) was
20 performed. This type of analysis allows for the synthesis of data from direct and indirect
21 comparisons without breaking randomisation and allows for the ranking of different
22 interventions. In this case the outcomes were defined as:

- 23 • probability of response.

24 These estimates provide a useful clinical summary of the results that can feed into an
25 economic model and facilitate the formation of recommendations based on the best available
26 evidence.

27 Conventional fixed effects meta-analysis assumes that the relative effect of one treatment
28 compared to another is the same across an entire set of trials. In a random effects model, it
29 is assumed that the relative effects are different in each trial but that they are from a single
30 common distribution and that this distribution is common across all sets of trials.

31 Network meta-analysis requires an additional assumption over conventional meta-analysis.
32 The additional assumption is that intervention A has the same effect on people in trials of
33 intervention A compared to intervention B as it does for people in trials of intervention A
34 versus intervention C, and so on. Thus, in a random effects network meta-analysis, the
35 assumption is that intervention A has the same effect distribution across trials of A versus B,
36 A versus C, and so on.

37 This specific method is usually referred to as mixed-treatment comparisons analysis but we
38 will continue to use the term network meta-analysis to refer generically to this kind of
39 analysis. We do so since the term “network” better describes the data structure, whereas
40 “mixed treatments” could easily be misinterpreted as referring to combinations of treatments.

41

1 1.2 Methods

2 1.2.1 Study selection

3 To estimate the probability of response, we performed an NMA that simultaneously used all
4 the relevant RCT evidence from the clinical evidence review. As with conventional meta-
5 analyses, this type of analysis does not break the randomisation of the evidence, nor does it
6 make any assumptions about adding the effects of different interventions. The effectiveness
7 of a particular intervention that involves a combination of interventions will be derived only
8 from randomised controlled trials that had that particular combination in a trial arm.

9 1.2.2 Outcome measures

10 The guideline clinical evidence reviews considered continuous outcomes to be the priority
11 outcomes that the committee wished to make recommendations from. Dichotomous
12 outcomes were also included but were only extracted from a study into the clinical review if
13 the study didn't report any continuous outcomes. Response to an intervention was the only
14 way to link outcomes to quality of life, and therefore only dichotomous outcomes could be
15 utilised in any modelling.

16 It would have been difficult to undertake an NMA for the guideline as a whole because of
17 issues with the populations in the studies being dissimilar in terms of medication status, and
18 the differences in the interventions being provided which would have been a particular issue
19 for the non-pharmacological and combination questions. And therefore seeing as the clinical
20 evidence used for the health economic model was already quite far removed from the larger
21 pool of evidence identified for the clinical review (both because of the small pool of evidence
22 being used for treatment effect in the model, and also because the outcomes being used in
23 the model are secondary outcomes on the clinical protocol), this NMA was undertaken by the
24 health economist purely to inform the economic model.

25 1.2.3 Comparability of interventions

26 The interventions compared in the model were a subset of those found in the randomised
27 controlled trials included in the clinical evidence review presented in evidence review F:
28 Combination treatment. Studies from the clinical review in the combination question first had
29 to be assessed for whether they reported dichotomous outcomes, which were needed for the
30 model. The comparisons in this pool of relevant studies were then extracted, and as
31 presented in section 1.2 of appendix 2, the rationale for what the comparators were in the
32 NMA were dependent on what comparisons were found in studies with dichotomous
33 outcomes, and whether the committee felt the studies could be pooled or not because of
34 similarity in interventions being assessed.

35 Treatments included in the network meta-analysis;

- 36 • Behavioural therapy
- 37 • Atomoxetine
- 38 • Combination of behavioural therapy and atomoxetine

39 1.2.4 Baseline risks

40 The baseline risk is defined as the risk of achieving the outcome of interest in the baseline
41 treatment arm of the included trials (i.e. the treatment labelled '1'). A meta-analysis was run
42 on the baseline separately to the NMA model.

1 1.2.5 Statistical analysis

2 A hierarchical Bayesian network meta-analysis (NMA) was performed using the software
3 WinBUGS⁴. We adapted fixed effects and random effects code from the NICE Decision
4 Support Unit¹ (by adding additional code to calculate the residual deviance for example). This
5 method accounts for the correlation between study level effects induced by multi-arm trials.

6 In order to be included in the analysis, a fundamental requirement is that each treatment is
7 connected directly or indirectly to every other intervention in the network. A diagram of the
8 evidence network is presented in section 1.3, with the detail of the comparisons in each
9 study in Table 1.

10 A baseline meta-analysis was undertaken as it is recommended by the NICE Decision
11 Support Unit that the same Generalised Linear Modelling framework is used to model the
12 absolute effects of a “standard treatment” or placebo comparator, as that proposed for
13 synthesis of relative treatment effects. This was informed by two studies, and the code can
14 be found in section 1.6. This used a fixed effects model instead of a random effects model,
15 as there weren’t enough studies to estimate the heterogeneity. However, the sample
16 probabilities of the two studies were similar.

17 Both the baseline meta-analysis and the NMA used a binomial likelihood logit link model.
18 Because we were interested in an outcome of the number of events (responses) out of the
19 total number of patients in each arm in each trial, we assumed that the data generation
20 process follows a Binomial likelihood. Since the parameters of interest were probabilities and
21 therefore can only take values between 0 and 1, a transformation (link function) was used
22 that maps these probabilities into a continuous measure between plus and minus infinity. For
23 a Binomial likelihood the most commonly used link function is the logit link function.

24 The NMA model used a fixed effects model, with parameters estimated by Markov chain
25 Monte Carlo simulation. Both a random effects and fixed effects model were tested and the
26 goodness of fit was compared. As there wasn’t much difference between the two, a fixed
27 effects model was used.

28 For the analyses, a series of 60,000 burn-in simulations were run to allow convergence and
29 then a further 60,000 simulations were run to produce the outputs. Convergence was
30 assessed by examining the history and kernel density plots.

31 We tested the goodness of fit of the model by calculating the residual deviance. If the
32 residual deviance is close to the number of unconstrained data points (the number of trial
33 arms in the analysis) then the model is explaining the data well.

34 The results, in terms of probability of response, are reported in section 1.3 below.

35 The aim of the NMA was to calculate the probability of response, for ease of interpretation,
36 and so that they could be easily fed into the economic model.

37 A key assumption behind an NMA is that the network is consistent. In other words, it is
38 assumed that the direct and indirect treatment effect estimates do not disagree with one
39 another. Discrepancies between direct and indirect estimates of effect may result from
40 several possible causes. First, there is chance and if this is the case then the network meta-
41 analysis results are likely to be more precise as they pool together more data than
42 conventional meta-analysis estimates alone. Second, there could be differences between the
43 trials included in terms of their clinical or methodological characteristics.

44 This heterogeneity is a problem for network meta-analysis but may be dealt with by subgroup
45 analysis, meta-regression or by carefully defining inclusion criteria. In this network it is
46 arguable if we need to check for consistency at all since the only loop is formed by a 3-arm
47 study which is consistent by design. However, we tested for inconsistency by fitting an
48 inconsistency model² for networks of binary outcomes. We compared the posterior mean of

the residual deviance between the consistency and inconsistency models to see which was a better fit to the data (closest to the number of trial arms in each network) and checked that the difference in deviance information criterion (DIC) values between the two models was small (less than 3-5), or if it was larger, that the smaller DIC and hence better fitting model was the consistency model. We also compared the direct and indirect evidence by testing if the (1,2) and (1,3) comparisons in the 3-arm trial, plus the (1,3) comparison in the 2-arm trial, agree with the (2,3) comparison in the other 2-arm trial. A p-value for inconsistency was also calculated. There was no evidence of inconsistency.

1.3 Results

1.3.1 Included studies

Three^{3,5,6} studies were identified as reporting dichotomous outcomes of response that also had relevant comparators (see more on this in the clinical data overview section of appendix 2 (section 1.2).

One study had 3 comparators³ and was the only study to form a closed loop, with the other two studies having 2 comparators^{5,6}.

The network can be seen in Figure 1, and the trial data for each of the studies included in the NMA are presented in

Table 1.

Figure 1: Network of studies used for treatment response

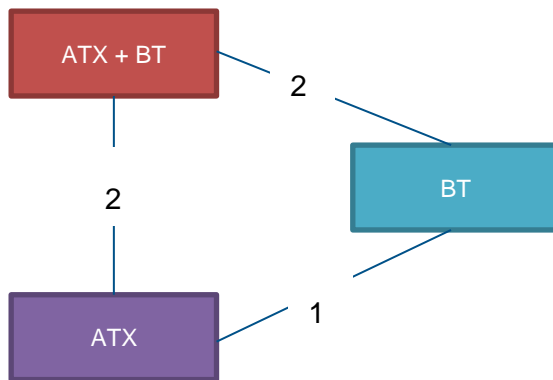


Table 1: Study data for ADHD network meta-analysis

Study	Intervention 1	Intervention 2	Intervention 3	Intervention 1		Intervention 2		Intervention 3	
				N	No. A	N	No. A	N	No. A
Handen 2015	Behavioural therapy	Atomoxetine	Combination	9	31	15	32	15	31
Waxmonsky 2010	Atomoxetine	Combination		14	27	16	29	-	-
Svanborg	Behavioural	Combination		14	50	35	49	-	-

Study	Intervention 1	Intervention 2	Intervention 3	Intervention 1		Intervention 2		Intervention 3	
				N	No. A	N	No. A	N	No. A
2009	therapy								

1 *N: number of events, No.A: number analysed*

2 Table 2 also summarises the studies in more detail.

3 **Table 2: Study detail**

Study	Intervention 1	Intervention 2	Intervention 3
Handen 2015 Aged 5-14. Mean age = around 8 in each group. 45.3% had received prior treatment for ADHD.	Behavioural therapy: (parent training plus placebo). Weekly meetings of 60-90 minutes. 9 meetings. A home visit was also conducted between the second and third session. N=31	Atomoxetine (ATX) "Final dose of 49.8mg or 1.3mg/kg. ATX doses were individually adjusted according to a weight-based dosage schedule, with medical clinicians allowed to delay increases or to reduce doses due to AEs. Initial dose = 0.3mg/kg/day (rounded to the nearest 5 mg) with weekly escalations by 0.3mg/kg/day, unless there were limiting side effects or no further room for improvement, to a target dose of 1.2 mg/kg/day, and could be increased to a maximum of 1.8 mg/kg/day based on clinical status and response" N=32	Combination Final dose of 40mg or 1.35mg/kg. Weekly 1:1 meetings of 60-90 minutes. Assumed for 10 weeks? A home visit was also conducted between the second and third session. N=31
Waxmonsky 2010 Aged 6-12. Mean 8.59. Some had previously taken ATX and some had started it before the trial. 37.5% had never taken stimulants. Excluded people who previously failed to respond to ATX.	Atomoxetine Medication provided in a single morning dose. Dose of 0.5mg/kg was started for 3 days then 0.8mg/kg for next 4 days, on day 8 everyone had dose increased to 1.2mg/kg. At 3 weeks tolerability was assessed and dose could be increased to 1.8mg/kg if CGI-S score was 4 or worse. Mean dose at study endpoint was 1.47mg/kg in ATX group. N=27	Combination Medication provided in a single morning dose. Dose of 0.5mg/kg was started for 3 days then 0.8mg/kg for next 4 days, on day 8 everyone had dose increased to 1.2mg/kg. At 3 weeks tolerability was assessed and dose could be increased to 1.8mg/kg if CGI-S score was 4 or worse. Mean dose at study endpoint was 1.40mg/kg in ATX+BT group. 3 components to BT; parenting program, social skills training, and school based daily report card. Sessions were weekly for 2 hrs in groups, children attended a simultaneous social skills program. N=29	
Svanborg 2009	Behavioural therapy	Combination	

Aged 6-15 Mean = 11.5 stimulant naïve children	(placebo + psycho-education) Parents participated in 4 session psycho-educational training. Four 3 hour group sessions. Contains components that might be more behavioural training. N=50	(ATX + psychoeducation) 0.5mg/kg during the first week, thereafter 1.2mg/kg (< or = 70 kg) or 80 mg/day (> 70 kg). It was dispensed at 6 visits (visits 2-7) during active treatment phase. Parents participated in 4 session psycho-educational training. Four 3 hour group sessions. Could be seen as more behavioural therapy as 'the content of the program contained core elements of more comprehensive behavioural treatment programs like parental management training (PMT) and the community parent education program (COPE)'. N=49	
--	---	--	--

1

2 In the NMA, behavioural therapy was chosen to be the baseline treatment that effects would
 3 be compared to. There is no 'placebo' comparison of those being compared as all are active
 4 treatments. Therefore behavioural therapy was chosen purely because it is the least effective
 5 from the data available, and if we had wanted to use ratios from the NMA output (such as
 6 odds ratios), then they would be more than 1 when compared to behavioural therapy. Current
 7 practice can also vary, and as the populations from the included trials were a mix of people
 8 on concurrent treatment or not then it is difficult to make an assumption about what the
 9 baseline would be (e.g. if it was newly diagnosed children with moderate impairment then
 10 using the last guideline as a guide; behavioural therapy might be the first line option and
 11 therefore the baseline). But as this was not the case because of the mixed population, the
 12 least effective was chosen.

13 **1.3.2 NMA results**

14 Table 3 summarises the results of the network meta-analysis in terms of the probability of
 15 response for each intervention, as well as the standard deviation, median, and confidence
 16 interval around each of the probabilities for each intervention.

17 **Table 3: NMA results: Posterior distribution of the probability of response for each**
 18 **intervention**

Intervention	Mean	SD	2.50% confidence interval	median	97.50% confidence interval
Behavioural therapy	0.2842	0.0501	0.1937	0.2814	0.3899
Atomoxetine	0.5666	0.1165	0.3327	0.5703	0.7810
Combination	0.6250	0.0950	0.4289	0.6296	0.7964

19 The probabilities of response were used directly in the model as the clinical outcomes.
 20 Quality of life was attached to the responders and non-responders, and also costs of the
 21 interventions were included to generate ICERS.

1 1.3.2.1 Goodness of fit and inconsistency

2 The fixed effects model used for the NMA was a good fit, with a residual deviance of 7.3
3 reported. This corresponds well to the total number of trial arms, 7.

4 An inconsistency model was run and the Deviance Information Criterion (DIC) statistics were
5 as follows in Table 4. The difference in the DIC is small (<3-5) with the consistency model
6 having the lower DIC value. This suggests that it fits the data better than the inconsistency
7 model.

8 A p-value for inconsistency was also calculated (0.78).

9 **Table 4: Goodness of fit statistics for the network meta-analysis and inconsistency**
10 **models**

	Deviance Information Criterion (DIC)	Posterior mean of the residual deviance (resdev)
Consistency model (fixed effects)	39.818	7.319
Inconsistency model (fixed effects)	41.746	8.256

11

12 1.4 Discussion

13 Based on the results of conventional meta-analyses of direct evidence, as has been
14 previously presented in evidence review F on combination treatment, deciding upon the most
15 clinical and cost effective intervention in an ADHD population is challenging. In order to
16 overcome the difficulty of interpreting the conclusions from numerous separate comparisons,
17 network meta-analysis of the direct evidence was performed. The findings of the NMA were
18 used to facilitate the guideline committee in decision-making when developing
19 recommendations.

20 The outcome chosen for the NMA was heavily influenced by the outcome being needed for
21 the health economic model. As previously explained, an NMA for the review as a whole
22 would have been difficult to undertake because of issues with the populations in the studies,
23 and the differences in the interventions being provided. This NMA was undertaken by the
24 health economist purely to inform the economic model.

25 Our analysis was based on a singular outcome or probability of response. 3 studies informed
26 the ADHD network where 3 different individual or combination treatments were evaluated
27 including a non-pharmacological treatment (behavioural therapy), a pharmacological
28 treatment (atomoxetine), and a combined non-pharmacological and pharmacological
29 intervention (atomoxetine + behavioural therapy).

30 The NMA was only informed by 3 studies, and therefore was a very small network. However
31 the alternatives would have been either; using the data in the model by crudely working out
32 the response rates through summation across the studies – which would break
33 randomisation. Or choosing one intervention as baseline and applying the relative effect of
34 the others – which can lead to different results depending on which arm was chosen as the
35 baseline, because the different direct comparisons are not identical. Therefore an NMA was
36 the most statistically robust way of combining the data. In the ADHD network, the intervention
37 with the highest probability of response was combination treatment, closely followed by
38 Atomoxetine, and finally behavioural therapy was the least effective (see Table 3). There was
39 a lot of uncertainty about the estimates with the credible intervals for some of the
40 interventions being very wide.

1 The network seemed to fit well, as demonstrated by DIC and residual deviance statistics.
2 However due to the limited number of studies, the credible intervals around the ranking of
3 treatments in the network was wide, suggesting considerable uncertainty about these results.

4 1.5 Conclusion

5 This analysis allowed us to combine findings from different comparisons presented in the
6 review even when direct comparative data was lacking.

7 For details of the rationale and discussion leading to recommendations, please refer to the
8 section 'the committee's discussion of the evidence' (section 1.11 of evidence review F on
9 combination treatment).

10 1.6 WinBUGS codes

11 1.6.1 Winbugs code for the baseline meta-analysis

```
12  
13 Atomoxetine baseline Data (BT arm)  
14 =====  
15 2 trials  
16  
17 =====  
18  
19 # Binomial likelihood, logit link  
20 # Baseline fixed effects model  
21 model{                                     # *** PROGRAM STARTS  
22 for (i in 1:ns){                           # LOOP THROUGH STUDIES  
23     r[i] ~ dbin(p[i],n[i])                 # Likelihood  
24     logit(p[i]) <- m                       # Log-odds of response  
25 #Deviance contribution  
26     rhat[i] <- p[i] * n[i] # expected value of the numerators  
27     dev[i] <- 2 * (r[i] * (log(r[i])-log(rhat[i])))  
28         + (n[i]-r[i]) * (log(n[i]-r[i]) - log(n[i]-rhat[i])))  
29     }  
30 totresdev <- sum(dev[])                    # Total Residual Deviance  
31 m ~ dnorm(0,.0001)                        # vague prior for mean  
32 logit(R) <- m                             # posterior probability of response  
33 }  
34  
35 Data  
36  
37 list(ns=2) # ns=number of studies  
38  
39 r[]    n[]    #    Study ID  
40 9      31    #    1  
41 14     50    #    3  
42  
43 END  
44  
45  
46  
47 Inits  
48 list(m=0)  
49  
50 list(m= -1)
```

1
2 list(m = 1)

3

4 1.6.2 Winbugs code for the probability of response

5

6 **ATX model**

7 **treatment 1 = BT**

8 **treatment 2 = ATX**

9 **treatment 3 = combo**

10

11 This code is part of

12 Dias, S., Welton, N.J., Sutton, A.J. & Ades, A.E. NICE DSU Technical Support Document 2: A Generalised Linear Modelling
13 Framework for Pairwise and Network Meta-Analysis of Randomised Controlled Trials. 2011; last updated September 2016
14 (available from <http://www.nicedsu.org.uk>).

15 This work should be cited whenever the code is used whether in its standard form or adapted.

16

17 # Binomial likelihood, logit link

18 # Fixed effects model

19 model{

20 for(i in 1:ns){ # *** PROGRAM STARTS

21 # LOOP THROUGH STUDIES

22 mu[i] ~ dnorm(0, .0001) # vague priors for all trial baselines

23 for (k in 1:na[i]) { # LOOP THROUGH ARMS

24 r[i,k] ~ dbin(p[i,k],n[i,k]) # binomial likelihood

25 # model for linear predictor

26 logit(p[i,k]) <- mu[i] + d[t[i,k]] - d[t[i,1]]

27 # expected value of the numerators

28 rhat[i,k] <- p[i,k] * n[i,k]

29 #Deviance contribution

30 dev[i,k] <- 2 * (r[i,k] * (log(r[i,k])-log(rhat[i,k])))

31 + (n[i,k]-r[i,k]) * (log(n[i,k]-r[i,k]) - log(n[i,k]-

32 rhat[i,k])))

33 }

34 # summed residual deviance contribution for this trial

35 resdev[i] <- sum(dev[i,1:na[i]])

36 }

37 totresdev <- sum(resdev[]) # Total Residual Deviance

38 d[1]<-0 # treatment effect is zero for reference treatment

39 # vague priors for treatment effects

40 for (k in 2:nt){ d[k] ~ dnorm(0, .0001) }

41 # obtain all pairwise ORs

42 for (c in 1:(nt-1)){

43 for (k in (c+1):nt) {

44 OR[c,k] <- exp(d[k] - d[c])

45 LOR[c,k]<-(d[k]-d[c])

46 }

47 }

48 # Provide estimates of treatment effects T[k] on the natural (probability)
49 scale

50 # Given a Mean Effect, meanA, for 'standard' treatment A,

51 # with precision (1/variance) precA

52 A ~ dnorm(meanA,precA)

53 for (k in 1:nt) { logit(T[k]) <- A + d[k] }

54 }

55 # *** PROGRAM ENDS

56

57 **Data**

58 # ns= number of studies; nt=number of treatments

list(ns=3, nt=3, meanA=-0.9378, precA=16.11582508)

```

1
2
3 r[,1] n[,1] r[,2] n[,2] r[,3] n[,3] t[,1] t[,2] t[,3] na[]
4 9 31 15 32 15 31 1 2 3 3
5 14 27 16 29 NA NA 2 3 NA 2
6 14 50 35 49 NA NA 1 3 NA 2
7
8
9
10
11
12
13
14
15
16
    
```

Initial Values

```

17 #chain 1
18 list(d=c( NA, 0, 0), mu=c(0, 0, 0))
19 #chain 2
20 list(d=c( NA, -1, 0), mu=c(-3, -3, -3))
21 #chain 3
22 list(d=c( NA, 2, 0), mu=c(-3, 4, 1))
23
24
25
    
```

1.6.3 Winbugs code for NMA inconsistency model

FE model for ATX model data: 3 trials, 3 treatments

```

26
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43
44
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48
49
50
51
52
53
54
55
56
=====
1 BT
2 ATX
3 combo
=====

# Binomial likelihood, logit link, inconsistency model
# Fixed effects model
model{
    # *** PROGRAM STARTS
    for(i in 1:ns){
        # LOOP THROUGH STUDIES
        mu[i] ~ dnorm(0,.0001) # vague priors for trial baselines
        for (k in 1:na[i]) { # LOOP THROUGH ARMS
            r[i,k] ~ dbin(p[i,k],n[i,k]) # binomial likelihood
            logit(p[i,k]) <- mu[i] + d[t[i,1],t[i,k]] # model for linear
        }
        predictor
        #Deviance contribution
        rhat[i,k] <- p[i,k] * n[i,k] # expected value of the numerators
        dev[i,k] <- 2 * (r[i,k] * (log(r[i,k])-log(rhat[i,k])))
            + (n[i,k]-r[i,k]) * (log(n[i,k]-r[i,k]) - log(n[i,k]-
            rhat[i,k])))
    }
    # summed residual deviance contribution for this trial
    resdev[i] <- sum(dev[i,1:na[i]])
}
totresdev <- sum(resdev[]) # Total Residual Deviance
for (k in 1:nt) { d[k,k] <- 0 } # set effects of k vs k to zero
for (c in 1:(nt-1)) { # priors for all mean treatment effects
    for (k in (c+1):nt) { d[c,k] ~ dnorm(0,.0001) }
}
# calculate prob of inconsistency
d.23 <- d[1,3]-d[1,2]
diff.23 <- d.23 - d[2,3]
prob <- step(diff.23)
} # *** PROGRAM ENDS
    
```

Data

```

1 # ns= number of studies; nt=number of treatments
2 list(ns=3, nt=3)
3
4 r[,1]    n[,1]    r[,2]    n[,2]    r[,3]    n[,3]    t[,1]    t[,2]    t[,3]    na[]
5 9        31       15       32       15       31       1        2        3        3
6 14       27       16       29       NA       NA       2        3        NA       2
7 14       50       35       49       NA       NA       1        3        NA       2
8
9 END
10
11
12 Inits
13 #chain 1
14 list(mu=c(-2,0,2),
15 d = structure(.Data = c(
16     NA,0,0,
17     NA,NA,0,
18     NA, NA, NA),
19 .Dim = c(3,3)) )
20
21 #chain 2
22 list(mu=c(1,0,3),
23 d = structure(.Data = c(
24     NA,2,2,
25     NA,NA,2,
26     NA, NA, NA),
27 .Dim = c(3,3)) )
28
29 #chain 3
30 list(mu=c(2,-2,1),
31 d = structure(.Data = c(
32     NA,1,1,
33     NA,NA,1,
34     NA, NA, NA),
35 .Dim = c(3,3)) )
36
37
38
39
    
```

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1 Appendices

2 Appendix A: Search strategy

3 4 A.1 Clinical search literature search strategy

5 Searches for patient views were run in Medline (OVID), Embase (OVID), CINAHL, Current
 6 Nursing and Allied Health Literature (EBSCO) and PsycINFO (ProQuest). Search filters were
 7 applied to the search where appropriate.

8 **Table 5: Database date parameters and filters used**

Database	Dates searched	Search filter used
Medline (OVID)	1948 – 28 April 2017	Exclusions Patient views/qualitative studies
Embase (OVID)	1974– 28 April 2017	Exclusions Patient views/qualitative studies
CINAHL (EBSCO)	Inception– 28 April 2017	Exclusions Patient views/qualitative studies
PsycINFO (ProQuest)	Inception– 28 April 2017	Exclusions Patient views/qualitative studies

9 **Medline (Ovid) search terms**

1.	"attention deficit and disruptive behavior disorders"/ or attention deficit disorder with hyperactivity/
2.	((attenti* or disrupt*) adj3 (adolescent* or adult* or behav* or child* or class or classes or classroom* or condition* or difficult* or disorder* or learn* or people or person* or poor or problem* or process* or youngster*).ti.
3.	((attenti* or disrupt*) adj3 disorder*).ab.
4.	(adhd or addh or ad hd or ad??hd).ti,ab.
5.	(attenti* adj3 deficit*).ti,ab.
6.	((hyperkin* or hyper kin*) adj1 (syndrome* or disorder*)) or hkd).ti,ab.
7.	(minimal brain adj2 (dysfunct* or disorder*).ti,ab.
8.	or/1-7
9.	limit 8 to English language
10.	letter/
11.	editorial/
12.	news/
13.	exp historical article/
14.	Anecdotes as Topic/
15.	comment/
16.	case report/
17.	(letter or comment*).ti.
18.	or/10-17

19.	randomized controlled trial/ or random*.ti,ab.
20.	18 not 19
21.	animals/ not humans/
22.	Animals, Laboratory/
23.	exp animal experiment/
24.	exp animal model/
25.	exp Rodentia/
26.	(rat or rats or mouse or mice).ti.
27.	or/20-26
28.	9 not 27
29.	Qualitative research/ or Narration/ or exp Interviews as Topic/ or exp "Surveys and Questionnaires"/ or Health care surveys/
30.	(qualitative or interview* or focus group* or theme* or questionnaire* or survey*).ti,ab.
31.	(metasynthes* or meta-synthes* or metasummar* or meta-summar* or metastud* or meta-stud* or metathem* or meta-them* or ethno* or emic or etic or phenomenolog* or grounded theory or constant compar* or (thematic* adj3 analys*) or theoretical sampl* or purposive sampl* or hermeneutic* or heidegger* or husserl* or colaizzi* or van kaam* or van manen* or giorgi* or glaser* or strauss* or ricoeur* or spiegelberg* or merleau*).ti,ab.
32.	or/29-31
33.	28 and 32

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Embase (Ovid) search terms

1.	attention deficit disorder/
2.	((attenti* or disrupt*) adj3 (adolescent* or adult* or behav* or child* or class or classes or classroom* or condition* or difficult* or disorder* or learn* or people or person* or poor or problem* or process* or youngster*).ti.
3.	((attenti* or disrupt*) adj3 disorder*).ab.
4.	(adhd or addh or ad hd or ad??hd).ti,ab.
5.	(attenti* adj3 deficit*).ti,ab.
6.	((hyperkin* or hyper kin*) adj1 (syndrome* or disorder*)) or hkd).ti,ab.
7.	(minimal brain adj2 (dysfunct* or disorder*).ti,ab.
8.	or/1-7
9.	limit 8 to English language
10.	letter.pt. or letter/
11.	note.pt.
12.	editorial.pt.
13.	case report/ or case study/
14.	(letter or comment*).ti.
15.	or/10-14
16.	randomized controlled trial/ or random*.ti,ab.
17.	15 not 16
18.	animal/ not human/
19.	nonhuman/
20.	exp Animal Experiment/
21.	exp Experimental Animal/

22.	animal model/
23.	exp Rodent/
24.	(rat or rats or mouse or mice).ti.
25.	or/17-24
26.	9 not 25
27.	health survey/ or exp questionnaire/ or exp interview/ or qualitative research/ or narrative/
28.	(qualitative or interview* or focus group* or theme* or questionnaire* or survey*).ti,ab.
29.	(metasynthes* or meta-synthes* or metasummar* or meta-summar* or metastud* or meta-stud* or metathem* or meta-them* or ethno* or emic or etic or phenomenolog* or grounded theory or constant compar* or (thematic* adj3 analys*) or theoretical sampl* or purposive sampl* or hermeneutic* or heidegger* or husserl* or colaizzi* or van kaam* or van manen* or giorgi* or glaser* or strauss* or ricoeur* or spiegelberg* or merleau*).ti,ab.
30.	or/27-29
31.	26 and 30

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CINAHL (EBSCO) search terms

S1.	(MH "Attention Deficit Hyperactivity Disorder")
S2.	((attenti* or disrupt*) n3 (adolescent* or adult* or behav* or child* or class or classes or classroom* or condition* or difficult* or disorder* or learn* or people or person* or poor or problem* or process* or youngster*))
S3.	adhd or addh or ad hd or ad/hd
S4.	attenti* n3 deficit*
S5.	((hyperkin* or hyper kin*) n1 (syndrome* or disorder*)) or hkd)
S6.	(minimal brain n2 (dysfunct* or disorder*))
S7.	S1 OR S2 OR S3 OR S4 OR S5 OR S6
S8.	(MH "Qualitative Studies+")
S9.	(MH "Qualitative Validity+")
S10.	(MH "Interviews+") OR (MH "Focus Groups") OR (MH "Surveys") OR (MH "Questionnaires+")
S11.	(qualitative or interview* or focus group* or theme* or questionnaire* or survey*)
S12.	(metasynthes* or meta-synthes* or metasummar* or meta-summar* or metastud* or meta-stud* or metathem* or meta-them* or ethno* or emic or etic or phenomenolog* or grounded theory or constant compar* or (thematic* n3 analys*) or theoretical sampl* or purposive sampl* or hermeneutic* or heidegger* or husserl* or colaizzi* or van kaam* or van manen* or giorgi* or glaser* or strauss* or ricoeur* or spiegelberg* or merleau*)
S13.	S8 OR S9 OR S10 OR S11 OR S12
S14.	S7 AND S13
S15.	PT anecdote or PT audiovisual or PT bibliography or PT biography or PT book or PT book review or PT brief item or PT cartoon or PT commentary or PT computer program or PT editorial or PT games or PT glossary or PT historical material or PT interview or PT letter or PT listservs or PT masters thesis or PT obituary or PT pamphlet or PT pamphlet chapter or PT pictorial or PT poetry or PT proceedings or PT "questions and answers" or PT response or PT software or PT teaching materials or PT website
S16.	S14 NOT S15 Limiters - English Language; Exclude MEDLINE records

3

4

PsycINFO (ProQuest) search terms

1.	SU.EXACT.EXPLODE("Attention Deficit Disorder") OR TI((attenti* OR disrupt*) NEAR/3 (adolescent* OR adult* OR behav* OR child* OR class OR classes OR classroom* OR condition* OR difficult* OR disorder* OR learn* OR people OR person* OR poor OR problem* OR process* OR youngster*)) OR AB((attenti* OR disrupt*) NEAR/3 disorder*) OR TI,AB(adhd OR addh OR ad-hd OR ad??hd) OR TI,AB(attenti* NEAR/3 deficit*) OR TI,AB(((hyperkin* OR (hyper-kin*)) NEAR/1 (syndrome* OR disorder*)) OR hkd) OR TI,AB(minimal NEAR/1 brain NEAR/2 (dysfunct* OR disorder*))
2.	SU.EXACT("Qualitative Research") OR (SU.EXACT("Narratives") OR SU.EXACT("Interviews")) OR (SU.EXACT("Questionnaires") OR SU.EXACT.EXPLODE("Surveys")) OR (qualitative OR interview*) OR (focus-group* OR theme*) OR (questionnaire* OR survey*) OR (metasynthes* OR meta-synthes*) OR (metasummar* OR meta-summar*) OR (metastud* OR meta-stud*) OR (metathem* OR meta-them*) OR ethno* OR (emic OR etic) OR (phenomenolog* OR "grounded theory") OR (constant-compar* OR thematic* NEAR/3 analys*) OR (theoretical-sampl* OR purposive-sampl*) OR (hermeneutic* OR heidegger*) OR (husserl* OR colaizzi*) OR (van-kaam* OR van-manen*) OR (giorgi* OR glaser*) OR (straus* OR ricoeur*) OR (spiegelberg* OR merleau*)
3.	1 AND 2
4.	NOT (Dissertations & Theses AND Books)
5.	English