# Bayesian Reasoning and Learning

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## 1 Introduction

Depression affects emotional, cognitive, and social well-being and manifests as prolonged low mood, diminished interest, hopelessness, fatigue, and altered sleep or appetite. Globally, around 280 million people suffer from depression World Health Organization [2023].

Conventional treatments include psychotherapy and pharmacological interventions Institute for Quality and Efficiency in Health Care [2024]. When these prove insufficient, alternative approaches like AI-driven mental healthcare offer significant promise by enabling early detection, personalized interventions, and virtual care. AI chatbots, in particular, enhance patient engagement and adherence Olawade et al. [2024], Liu et al. [2022].

Recent studies identify four dimensions of depression risk factors—socioeconomic status, physical attributes, genetic predispositions, and adverse childhood experiences Dagnino et al. [2020]. This study uses Bayesian network modeling to explore the interplay among these factors and depression onset, extending the framework with interventional nodes for both AI-based and traditional treatments to assess their moderating effects.

# 2 Domain

Depression stems from multiple interacting factors. For clarity, this study limits predictors to a select few. In the sociodemographic domain, only education and income are considered Dagnino et al. [2020]. In the physical domain, the focus is on physical activity and obesity status—highlighting obesity's impact over traditional BMI measurements (World Health Organization [2021b,a], Rutherford et al. [2022]). Genetic predisposition is modeled multiplicatively, reflecting that a family history can more than double depression risk (Dunn et al. [2015], Smith [2014]). Adverse childhood experiences (ACE) is incorporated using approximate estimates to maintain parsimony (Martins-Monteverde et al. [2019], Swedo et al. [2023]). Finally, nodes for AI-driven interventions and traditional treatments act as dampening factors in mitigating depression risk. This framework motivates the following research questions:

**Research Question 1**: How do sociodemographic factors, physical attributes, genetic influences, and adverse childhood experiences interact within a Bayesian network to predict depressive outcomes?

Research Question 2: How do traditional support methods and AI-enabled interventions interact to influence depression risk trajectories and outcomes within a Bayesian network framework?

# 3 Model

We use Python and the PyAgrum framework pyAgrum 2.1.1 Documentation [2025] to build our Bayesian network. Each variable represents a distinct determinant of depression, allowing us to examine their interrelationships. Table 1 lists the key variables in our simplified model. Nodes and arcs are then added using PyAgrum, as illustrated in the following code example.

# Create the combined Bayesian network for the depression model
bn = gum.BayesNet("CombinedDepressionModel")

Variable	Domain	Description
Education	{Low, Medium, High}	Low corresponds to primary education, VMBO, the lower sec-
		ondary phase of Havo/VWO, and MBO1. Medium denotes
		the upper secondary phase of Havo/VWO, MBO2, MBO3,
		and MBO4. High corresponds to HBO/WO. Data are ob-
		tained from Centraal Bureau voor de Statistiek [2022].
Income	{Low, Medium, High}	Categorizes household income in the USA, with data sourced
		from Pew Research Center [2020].
Physical Activity	{Inactive, Active}	Reflects the global percentage of the population that is phys-
		ically inactive [World Health Organization, 2021a].
Obesity	{Not Obese, Obese}	Represents the proportion of the global population classified
		as obese [World Health Organization, 2021b].
Socioeconomic Status (SS)	{High, Low}	Synthesizes key indicators of High or Low economic status.
, ,		Outcomes computed through a Noisy-OR.
Physical Attribute (PA)	{Poor, Good}	Synthesizes key indicators of physical health. Data is esti-
		mated heuristically for educational purposes
Adverse Childhood Experience (ACE)	{Absent, Present}	Indicates whether adverse childhood experiences were encoun-
		tered. Data are retrieved from Swedo et al. [2023].
Genetic Predisposition (GP)	{Absent, Present}	Denotes the presence or absence of a genetic risk factor for
		depression. Data are retrieved from Kobayashi et al. [2024].
Depression	{No, Yes}	Binary outcome representing the absence or presence of de-
		pression. Outcomes computed through an Adapted Leaky
		Noisy-OR.
Artificial Intelligence	{No, Yes}	A binary variable denoting whether artificial intelligence has
		been implemented as an intervention to mitigate depression.
		The impact is estimated heuristically for educational pur-
		poses.
Traditional Methods	{No, Yes}	A binary variable denoting whether conventional treatment
		modalities have been employed to manage depression. The im-
		pact is estimated heuristically for educational purposes.

Table 1: Key variables in the simplified depression model.

Determinants of depression are grouped into four domains. The SocioStatus variable is computed using a Noisy-OR gate and then combined with other factors via an adapted leaky Noisy-OR mechanism to yield the final outcomes. Figure 1 shows the resulting structure, and the code used to generate this graph is provided below.

```
node_colors = {bn.variable(i).name(): 1 for i in range(bn.size())}

node_colors["Artificial_Intelligence"] = 0.1
node_colors["Traditional_Methods"] = 0.1
node_colors["SocioStatus"] = 0.2
node_colors["Depression"] = 0.3
graph = gnb.show(bn, size="11", nodeColor=node_colors)
```

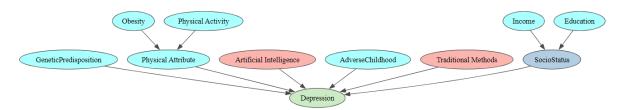


Figure 1: Bayesian network: red nodes are inactive by default, dark blue nodes use a standard Noisy-OR, and the green node uses a Adapted Leaky Noisy-OR.

#### 3.1 Noisy-OR

In the Noisy-OR model, the probability that a child variable X remains inactive  $(x^0)$  given its parents U is computed by

$$P(x^0 \mid U) = \prod_{u_i \in U} (1 - \lambda_i),$$

where  $\lambda_i$  is the probability that the *i*-th parent (here, SocioStatus) activates X. Thus, the probability that X is active  $(x^1)$  is

$$P(x^1 \mid U) = 1 - \prod_{u_i \in U} (1 - \lambda_i).$$

Table 2 lists the probabilities used for the SocioStatus variable.

Parameter	Key	State/Interpretation	Value
	0	Low	0.8
$\lambda_{Income}$	1	Medium	0.5
	2	High	0.2
	0	Low	0.7
$\lambda_{Education}$	1	Medium	0.4
	2	High	0.1

Table 2: Parameters used in the Noisy-OR for SocioStatus

Code used to compute SocioStatus:

```
cpt_socio = bn.cpt(socio)

r_inc = {"Low": 0.8, "Medium": 0.5, "High": 0.2}
r_edu = {"Low": 0.7, "Medium": 0.4, "High": 0.1}

income_states = bn.variable(inc).labels() # e.g., ["Low", "Medium", "High"]
edu_states = bn.variable(edu).labels() # e.g., ["Low", "Medium", "High"]

for inc_state, edu_state in itertools.product(income_states, edu_states):
    factor_inc = 1 - r_inc[inc_state]
    factor_edu = 1 - r_edu[edu_state]

product = factor_inc * factor_edu

p_low = 1 - product
p_high = product # so that p_high + p_low = 1

cpt_socio[{
    bn.variable(inc).name(): inc_state,
    bn.variable(edu).name(): edu_state,
```

# 3.2 Adapted Leaky Noisy-OR

When nodes for genetic predisposition, traditional methods, and AI-driven interventions are active, the model adjusts by computing  $P(x^1 \mid U)$  with depression-specific parameters. We define the adjusted probability as

$$P(\tilde{x}^1 \mid U) = \min \left\{ 1, \prod_{j \in M} r_j \left( P(x^1 \mid U) \right) \right\}. \tag{1}$$

In this equation,  $\lambda_i$  (see Table 3) represents the direct influences, and the  $r_j$  factors (Table 4) serve as multiplicative scaling modifiers. A latent baseline risk,  $\lambda_{\rm Leak}$ , accounts for depression risk from factors not explicitly modeled. All risk configurations are evaluated, with outcomes detailed in Tables 5–8 in the Probabilities subsection.

Parameter	Key	State/Interpretation	Value
1	0	Favorable (High SES)	0.005
$\lambda_{SS}$	1	Unfavorable (Low SES)	0.05
1	0	Favorable (Good health)	0.005
$\lambda_{PA}$	1	Unfavorable (Poor health)	0.03
1	0	Absent	0.005
$\lambda_{ACE}$	1	Present	0.03
$\lambda_{\rm Leak}$ – baseli	ine risk	(always present)	0.01

Table 3: Parameters triggering the Adapted Noisy-OR for Depression Outcome

Intervention	Condition	$r_j$
Genetic Predisposition	Present (doubles risk)	2
Artificial Intelligence	Present (reduces risk)	0.80
Traditional Methods	Present (reduces risk)	0.90
Combined Intervention	AI & Traditional Methods Present	0.70

Table 4: Effective Risk Multipliers

Code for computing the Depression node:

```
ss_params = \{0: 0.005, 1: 0.05\}
ph_params = \{0: 0.005, 1: 0.03\}
ace_params = \{0: 0.005, 1: 0.03\}
gp_multiplier = {0: 1.0, 1: 2}
chatbot = \{0: 1.0, 1: 0.80\}
traditional = \{0: 1.0, 1: 0.9\}
leak = 0.01
cpt_depress = bn.cpt(depress)
cpt_values = [] # This flat list will accumulate the CPT probabilities for all
    \hookrightarrow combinations
for th, cb, ace_state, gp_state, ph, ss in itertools.product(range(2), range(2),
    \hookrightarrow range(2), range(2), range(2), range(2)):
   prod_failure = (1 - ss_params[ss]) * (1 - ph_params[ph]) * (1 - ace_params[
        → ace_state]) * (1 - leak)
   p_baseline = 1 - prod_failure
   if gp_state == 1:
       p_dep = min(1.0, p_baseline * gp_multiplier[1])
    else:
```

```
p_dep = p_baseline
if cb == 1 and th == 1:
    p_dep = p_dep * 0.70
elif th == 1:
    p_dep = p_dep * traditional[1]
elif cb == 1:
    p_dep = p_dep * chatbot[1]
cpt_values.extend([1 - p_dep, p_dep])
cpt_depress.fillWith(cpt_values)
```

# 4 Results

#### 4.1 Probabilities

Data from the sources listed in Table 1 were cleaned and processed to derive the corresponding probability estimates. Tables 5 to 8 summarize depression outcomes under nonintervention and intervention scenarios. Due to the wide scope of the Depression variable, its representation was split into several smaller, manageable tables. Figures 2 through 9 depict the probability distributions for key variables including education, income, obesity, genetic predisposition, physical activity, physical attribute, adverse childhood experiences, and socioeconomic status. The code for generating these visualizations is provided below.

```
# Iterate over all nodes in the BN and show their CPTs:
for node in bn.nodes():
   var_name = bn.variable(node).name()
   print(f"Statistics_for_{var_name}:")
   gnb.showPotential(bn.cpt(node))
```

Adverse Childhood	Genetic Predisposition	Physical Attribute	SocioStatus	Depression (No)	Depression (Yes)
Absent	Absent	Poor	High	0.9752	0.0248
Absent	Absent	Poor	Low	0.9311	0.0689
Absent	Absent	Good	High	0.9507	0.0493
Absent	Absent	Good	Low	0.9077	0.0923
Absent	Present	Poor	High	0.9504	0.0496
Absent	Present	Poor	Low	0.8622	0.1378
Absent	Present	Good	High	0.9014	0.0986
Absent	Present	Good	Low	0.8154	0.1846
Present	Absent	Poor	High	0.9507	0.0493
Present	Absent	Poor	Low	0.9077	0.0923
Present	Absent	Good	High	0.9268	0.0732
Present	Absent	Good	Low	0.8849	0.1151
Present	Present	Poor	High	0.9014	0.0986
Present	Present	Poor	Low	0.8154	0.1846
Present	Present	Good	High	0.8537	0.1463
Present	Present	Good	Low	0.7698	0.2302

Table 5: Depression probabilities in default case

Adverse Childhood	Genetic Predisposition	Physical Attribute	SocioStatus	Depression (No)	Depression (Yes)
Absent	Absent	Good	High	0.9802	0.0198
Absent	Absent	Good	Low	0.9449	0.0551
Absent	Absent	Poor	High	0.9606	0.0394
Absent	Absent	Poor	Low	0.9262	0.0738
Absent	Present	Good	High	0.9604	0.0396
Absent	Present	Good	Low	0.8898	0.1102
Absent	Present	Poor	High	0.9212	0.0788
Absent	Present	Poor	Low	0.8524	0.1476
Present	Absent	Good	High	0.9606	0.0394
Present	Absent	Good	Low	0.9262	0.0738
Present	Absent	Poor	High	0.9415	0.0585
Present	Absent	Poor	Low	0.9079	0.0921
Present	Present	Good	High	0.9212	0.0788
Present	Present	Good	Low	0.8524	0.1476
Present	Present	Poor	High	0.8829	0.1171
Present	Present	Poor	Low	0.8159	0.1841

Table 6: Depression probabilities with Artificial Intelligence Intervention

Adverse Childhood	Genetic Predisposition	Physical Attribute	SocioStatus	Depression (No)	Depression (Yes)
Absent	Absent	Good	High	0.9777	0.0223
Absent	Absent	Good	Low	0.9380	0.0620
Absent	Absent	Poor	High	0.9556	0.0444
Absent	Absent	Poor	Low	0.9170	0.0830
Absent	Present	Good	High	0.9554	0.0446
Absent	Present	Good	Low	0.8760	0.1240
Absent	Present	Poor	High	0.9113	0.0887
Absent	Present	Poor	Low	0.8339	0.1661
Present	Absent	Good	High	0.9556	0.0444
Present	Absent	Good	Low	0.9170	0.0830
Present	Absent	Poor	High	0.9342	0.0658
Present	Absent	Poor	Low	0.8964	0.1036
Present	Present	Good	High	0.9113	0.0887
Present	Present	Good	Low	0.8339	0.1661
Present	Present	Poor	High	0.8683	0.1317
Present	Present	Poor	Low	0.7928	0.2072

Table 7: Depression probabilities with Traditional Methods Intervention

Adverse Childhood	Genetic Predisposition	Physical Attribute	SocioStatus	Depression (No)	Depression (Yes)
Absent	Absent	Good	High	0.9827	0.0173
Absent	Absent	Good	Low	0.9518	0.0482
Absent	Absent	Poor	High	0.9655	0.0345
Absent	Absent	Poor	Low	0.9354	0.0646
Absent	Present	Good	High	0.9653	0.0347
Absent	Present	Good	Low	0.9036	0.0964
Absent	Present	Poor	High	0.9310	0.0690
Absent	Present	Poor	Low	0.8708	0.1292
Present	Absent	Good	High	0.9655	0.0345
Present	Absent	Good	Low	0.9354	0.0646
Present	Absent	Poor	High	0.9488	0.0512
Present	Absent	Poor	Low	0.9194	0.0806
Present	Present	Good	High	0.9310	0.0690
Present	Present	Good	Low	0.8708	0.1292
Present	Present	Poor	High	0.8976	0.1024
Present	Present	Poor	Low	0.8389	0.1611

Table 8: Depression probabilities with both intervention methods

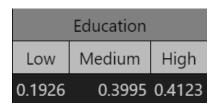


Figure 2: Education

Income					
Low Medium High					
0.0900	0.4300	0.4800			

Figure 3: Income

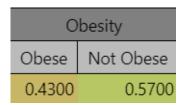


Figure 4: Obesity

GeneticPredisposition				
Absent Present				
0.6000 0.4000				

Figure 5: Genetic Predisposition

Physical Activity				
Inactive Active				
0.3100 0.6900				

Figure 6: Physical Activity

		Physical Attribute		
Obesity	Physical Activity	Good	Poor	
Obese	Inactive	0.9000	0.1000	
	Active	0.8000	0.2000	
N I O	Inactive	0.6000	0.4000	
Not Obese	Active	0.5000	0.5000	

Figure 7: Physical Attribute

AdverseChildhood		
Absent	Present	
0.3610	0.6390	

Figure 8: Adverse Childhood Experience



Figure 9: Socioeconomic Status

#### 4.2 Inference

The Bayesian network yields a baseline depression probability of 10.39% under default conditions. With the Artificial Intelligence intervention, this drops to 8.31%, while the Traditional Methods intervention lowers it to 9.35%. The combined intervention further reduces the probability to 7.27%. The following code computes and visualizes these outcomes:

```
gnb.showInference(bn, evs={})
gnb.showInference(bn, evs={"Artificial_Intelligence": "Yes"})
gnb.showInference(bn, evs={"Traditional_Methods": "Yes"})
gnb.showInference(bn, evs={
"Artificial_Intelligence": "Yes",
"Traditional_Methods": "Yes"
})
```

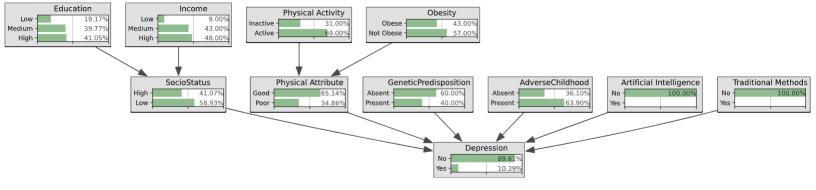
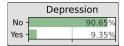


Figure 10: Default Outcome (10.39% probability)

Depression		
No -		91.69%
Yes -		8.31%



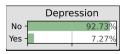


Figure 11: Artificial Intelligence Figure 12: Traditional Method Figure 13: Combined Outcome Outcome (8.31% probability)

Outcome (9.35% probability)

(7.27% probability)

#### 5 Discussion

For educational purposes, the model has been significantly simplified to highlight the essential mechanisms and interactions of risk factors within the Bayesian network, though it may not capture the full complexity of depression.

Data were drawn from diverse international sources, which may introduce discrepancies in risk factor measurements and interpretations due to variation in population characteristics and data collection

Additionally, some parameter values—particularly the trigger factors for the Noisy-ORs—are based on estimates to provide a tractable example rather than definitive clinical or epidemiological values.

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