1 Methodology supplement

Notation. We follow the notations in the main paper. Additionally, we use $E_{ij}$ to denote a matrix which has 1 at the $(i,j)$-th entry and 0 elsewhere.

1.1 An identifiable variable of CLG

We first show the copula correlation parameter $\Sigma$ in Definition 1 in the main paper is not identifiable.

**Theorem 1.1.** For a 2-dimensional categorical vector $(x_1, x_2) \sim \text{CLG}(\Sigma, \mu)$, $\text{CLG}(\Sigma^{(kl)}, \mu)$ has the same distribution as $\text{CLG}(\Sigma, \mu)$ where $\Sigma^{(kl)} = \Sigma_{[1],[2]} + c_1 \sum_{m=1}^K E_{km} + c_2 \sum_{m=1}^K E_{ml}$ for any constants $c_1, c_2$ and any integers $k, l = 1, \ldots, K$. In other words, adding any constant to a row or column in $\Sigma_{[1],[2]}$ does not change the distribution of $\text{CLG}(\Sigma, \mu)$.

**Proof.** For $(x_1, x_2) \sim \text{CLG}(\Sigma, \mu)$, denote the probability that $x_1 = i$ and $x_2 = j$ by $P_{ij}(\Sigma_{[1],[2]}, \mu)$ for $i, j = 1, \ldots, K$. To show $\text{CLG}(\Sigma, \mu)$ and $\text{CLG}(\Sigma^{(kl)}, \mu)$ have the same distribution, it suffices to show that $P_{ij}(\Sigma_{[1],[2]}, \mu) = P_{ij}(\Sigma^{(kl)}, \mu)$ for any $i, j = 1, \ldots, K$.

Now define $\Delta_i \in \mathbb{R}^{K-1 \times K}$ such that: (1) the $i$-th column of $\Delta_i$ has all entries equal to $-1$; (2) the remaining $K-1$ columns of $\Delta_i$ excluding the $i$-th column is $I_{K-1}$, the identity matrix. Then, we find that

$$i = \text{argmax}(z_i + \mu_{[1]}) \iff \Delta_i (z_i + \mu_{[1]}) \leq 0.$$

Further, define $\Delta_{i,j} = \text{diag}(\Delta_i, \Delta_j)$, then

$$x_1 = i, x_2 = j \iff \Delta_{i,j} z + \Delta_{i,j} \mu \leq 0.$$

Note $\Delta_{i,j} z$ is Gaussian distributed with mean zero and covariance matrix

$$\begin{bmatrix} \Delta_i \Delta_i^\top & \Delta_i \Sigma_{[1],[2]} \Delta_j^\top \\ \Delta_j \Sigma_{[2],[1]} \Delta_i^\top & \Delta_j \Delta_j^\top \end{bmatrix},$$

thus $P_{ij}(\Sigma_{[1],[2]}, \mu)$ depends on $\Sigma_{[1],[2]}, \mu$ only through $\Delta_i \Sigma_{[1],[2]} \Delta_j^\top$ and $\Delta_{i,j} \mu$. Now denote $R = \Sigma_{[1],[2]}$ and $R^{(kl)} = R + c_1 \sum_{m=1}^K E_{km} + c_2 \sum_{m=1}^K E_{ml}$ to simplify the notation. Note the $(s, t)$-th entry of $\Delta_i R \Delta_j^\top$ is $r_{ij} + r_{st} - r_{it} - r_{sj}$ and the $(s, t)$-th entry of $R^{(kl)}$ is $r_{st} + c_1 1(k = s) + c_2 1(l = t)$, then it is straightforward that $\Delta_i R \Delta_j^\top = \Delta_i R^{(kl)} \Delta_j^\top$ for arbitrary $c_1, c_2$ and any $k, l = 1, \ldots, K$, which finishes our proof.

To eliminate the unidentifiability showed in Theorem 1.1 we provide a variant of CLG with additional constraints in the copula correlation matrix $\Sigma$. Concretely, for each categorical variable $x_j$, we select one dimension in $z_{[j]}$ to be the base dimension, which does not correlate with any other entry in $z$. Without loss of generality, we can select the first dimension of $z_{[j]}$ for all $j$. In other words,
We prove the existence by contradiction. Suppose there is no satisfying $\lambda$ w.r.t. $e$. Also define $\Sigma$. The remaining computation is straightforward.

There are two motivations for this variant. First, as mentioned in Sec 2.1.2 of the main paper, to describe the joint distribution of two categorical variables (each has $K$ categories), $(K-1)^2$ free parameters are sufficient once given the marginal categorical distribution. Second, for a categorical variable $x$ generated as the argmax of a $K$-dim latent Gaussian $z$, only the difference among the entries of $z$ matters for the distribution of $x$. In fact, we can even fix the first dimension of $z$ to be constant 0 and Theorem 1 of the main paper still holds.

1.2 Computing the expectation of latent Gaussian

Here we show that computing $\mathbb{E}[zz^T|x_O, \Sigma]$ reduces to compute $\mathbb{E}[z_{[O]}|x_O, \Sigma]$ and $\text{Cov}[z_{[O]}|x_O, \Sigma]$. By writing $z = (z_{[O]}, z_{[M]})$, we first decompose the computation into three parts: (1) $\mathbb{E}[z_{[O]}z_{[O]^T}|x_O, \Sigma]$; (2) $\mathbb{E}[z_{[O]}z_{[M]^T}|x_O, \Sigma]$; (3) $\mathbb{E}[z_{[M]}z_{[M]^T}|x_O, \Sigma]$, and then show each of the three parts can reduce to the mean and covariance of $z_{[O]}|x_O, \Sigma$. For (1), it is trivial. For (2) and (3), the key technique is the law of total expectation and that

$$z_{[M]}|z_{[O]} \sim N \left( \Sigma_{[M],[O]}z_{[O]}, \Sigma_{[M],[O]} - \Sigma_{[M],[O]}\Sigma_{[O],[O]}^{-1}\Sigma_{[O],[M]} \right).$$

Thus

$$\mathbb{E}[z_{[O]}z_{[M]^T}|x_O, \Sigma] = \mathbb{E}_{z_{[O]}}[z_{[O]}\mathbb{E}_{z_{[M]|z_{[O]}}}[z_{[M]^T}|x_O, \Sigma]],$$

and

$$\mathbb{E}[z_{[M]}z_{[M]^T}|x_O, \Sigma] = \mathbb{E}_{z_{[M]}}[\mathbb{E}_{z_{[M]|z_{[O]}}}[z_{[M]}z_{[M]^T}|x_O, \Sigma]].$$

The remaining computation is straightforward.

1.3 Proof of Theorem 1

Proof. Denote $\mathbb{P}(\text{argmax}(z + \mu) = k) = p_k(\mu)$ for $k = 1, ..., K$ and $\mathbb{P}(\mu) = (p_1(\mu), ..., p_K(\mu))$. Also define $e_k$: $e_k \in \mathbb{R}^K$ has 1 at coordinate $k$ and zero elsewhere.

We prove the existence by contradiction. Suppose there is no satisfying $\mu$. Let

$$\mu^* = \arg\min_\mu f(\mu), \; \text{where} \; f(\mu) = \sum_{k=1}^K |p_k(\mu) - p_k|.$$  \hspace{1cm} (1)

Define $I = \{i | p_i(\mu^*) < p_i, i = 1, ..., K\}$ and $I^c = \{1, ..., K\} - I$. If there is no satisfying $\mu$, then both $I$ and $I^c$ are not empty. Pick an $i \in I$. Since $p_i(\mu^* + \lambda e_i)$ is a continuous function w.r.t. $\lambda$ and $\lim_{\lambda \to \infty} p_i(\mu^* + \lambda e_i) = 1$, there exists a $\lambda_0 > 0$ such that $p_i(\mu^* + \lambda_0 e_i) = p_i$. Note $p_k(\mu^* + \lambda e_i)$ is strictly decreasing w.r.t. $\lambda$ for any $k \neq i$. Thus for $p_k(\mu^* + \lambda_0 e_i) = p_k(\mu^*) - \delta_k$,
we have $\delta_k > 0$ when $k \neq i$ and $\delta_i = p_i(\mu^*) - p_i < 0$. Now

$$\sum_{k=1}^{K} |p_k(\mu^* + \lambda_0 \mathbf{e}_i) - p_k| = |p_i(\mu^* + \lambda_0 \mathbf{e}_i) - p_i| + \sum_{k \neq i} |p_k(\mu^* + \lambda_0 \mathbf{e}_i) - p_k|$$

$$= \sum_{k \in I - \{i\}} |p_k - p_k(\mu^*) + \delta_k| + \sum_{k \in I} |p_k - p_k(\mu^*) + \delta_k|$$

$$\leq \sum_{k \in I - \{i\}} (p_k - p_k(\mu^*) + \delta_k) + \sum_{k \in I} |p_k - p_k(\mu^*)| + \sum_{k \neq i} \delta_k$$

$$\leq \sum_{k \neq i} |p_k - p_k(\mu^*)| + \sum_{k \neq i} \delta_k$$

$$= f(\mu^*) - (p_i - p_i(\mu^*)) + \sum_{k \neq i} \delta_k$$

$$= f(\mu^*) + \sum_{k=1}^{K} \delta_k = f(\mu^*)$$

The equality only holds when for each $i \in I^\circ$, $p_k - p_k(\mu^*) \geq 0$, which further leads to $p_k = p_k(\mu^*)$ by the definition of $I^\circ$. That conflicts with that $I$ is nonempty. Thus we must have

$$f(\mu^* + \lambda_0 \mathbf{e}_i) < f(\mu^*),$$

which contradicts our assumption and completes our proof for existence.

Now for uniqueness, assume there exists $\mu \neq \hat{\mu}$ such that $P(\mu) = P(\hat{\mu})$. Define $I_\prec, I_\succ, I_\approx$ to be the set of entries that $\mu$ is smaller, equal, and larger than $\hat{\mu}$, respectively. We want to show it must be the case that both $I_\prec$ and $I_\succ$ are empty, which contradicts the assumption.

First, if $I_\prec$ is empty but $I_\succ$ is not, then for each $i \in I_\succ$, we define $\mu_i \in \mathbb{R}^K$ such that $\mu_i$ agrees with $\mu$ at all entries but $i$ and agrees with $\hat{\mu}$ at entry $i$. Since $p_i(\mu)$ is strictly increasing w.r.t. $\mu_i$ when fixing $\mu_{\{i\}}$, we know $p_i(\mu_i) < p_i(\mu)$. Further repeatedly switching one more entry of $\mu_i$ in $I_\succ$ from $\mu$ to $\hat{\mu}$ until $I_\succ$ is exhausted, we have $p_i(\mu_i) < p_i(\mu)$, which contradicts the assumption. Similarly we can show the contradiction if $I_\prec$ is not empty but $I_\succ$ is empty.

Now consider the case that both $I_\prec$ and $I_\succ$ are not empty. Define $\mu^*$ such that $\mu^*$ agrees with $\mu$ over $I_\prec$ and agrees with $\hat{\mu}$ over $I_\succ$. According to above, we know for each $i \in I_\prec$, $p_i(\mu) > p_i(\mu^*) > p_i(\hat{\mu})$, which contradicts the assumption. Thus we complete our proof. $\square$

2 Experiments supplement

2.1 Implementation details

All our implementation codes are provided in a Github repo[1]. All experiments use a laptop with a 3.1 GHz Intel Core i5 processor and 8 GB RAM. All algorithms are implemented using one core, although some of them including our EGC support parallelism.

The algorithm of EGC consists of two parts: the marginal estimation and the correlation estimation. The marginal estimation requires a subroutine to iteratively solve nonlinear systems. We find the available iterative root finding algorithms package in R such as the rootSolve do not achieve desired precision occasionally, while the scipy.optimize.root(method='hybr') function in Python finds accurate solution in all of our experiments. Through our experiments, EGC uses a Python implementation to estimate the marginal and a R implementation to estimate the copula correlation.

Table 1: Algorithm runtime in seconds: mean (sd) over 10 repetitions. The synthetic dataset is under the setting $K = 6, p_{cat} = 5$.

<table>
<thead>
<tr>
<th></th>
<th>EGC</th>
<th>missForest</th>
<th>MICE</th>
<th>ImputeFAMD</th>
<th>softImpute</th>
</tr>
</thead>
<tbody>
<tr>
<td>Synthetic</td>
<td>22.0 (1.0)</td>
<td>58.7 (9.9)</td>
<td>82.3 (1.0)</td>
<td>56.0 (22.3)</td>
<td>1.4 (0.2)</td>
</tr>
<tr>
<td>Abalone</td>
<td>9.8 (0.1)</td>
<td>136.2 (37.2)</td>
<td>2.9 (0.1)</td>
<td>46.8 (5.4)</td>
<td>0.9 (0.1)</td>
</tr>
<tr>
<td>Heart</td>
<td>2.1 (0.8)</td>
<td>2.6 (0.8)</td>
<td>4.3 (0.3)</td>
<td>11.7 (3.7)</td>
<td>0.1 (0.0)</td>
</tr>
<tr>
<td>CMC</td>
<td>5.5 (1.1)</td>
<td>9.2 (1.4)</td>
<td>17.3 (0.9)</td>
<td>41.2 (16.2)</td>
<td>0.3 (0.0)</td>
</tr>
<tr>
<td>Creditg</td>
<td>16.0 (0.9)</td>
<td>43.5 (12.3)</td>
<td>74.4 (1.2)</td>
<td>38.8 (11.3)</td>
<td>1.0 (0.1)</td>
</tr>
<tr>
<td>Credita</td>
<td>10.4 (1.1)</td>
<td>15.6 (3.2)</td>
<td>30.0 (8.6)</td>
<td>23.8 (12.1)</td>
<td>0.6 (0.1)</td>
</tr>
<tr>
<td>Colic</td>
<td>3.5 (0.2)</td>
<td>6.9 (2.0)</td>
<td>40.0 (15.3)</td>
<td>18.2 (15.8)</td>
<td>0.4 (0.1)</td>
</tr>
</tbody>
</table>

A complete R implementation is available though, and a complete Python implementation will be available soon. All other algorithms are completely implemented in R.

MICE is a multiple imputation method. To derive a single imputation from MICE, we pool 5 imputed datasets (majority vote for categorical and mean for ordered). We choose the rank for imputeFAMD in the grid of $\{1, 3, 5, 7, 9\}$. We choose the regularization parameter for softImpute in an exponentially decaying path of length 10 from $0.1 \times \lambda_0$ and $0.99 \times \lambda_0$, where $\lambda_0$ is computed using the provided function lambda0() in the package softImpute.

$$\exp(\text{seq}(\text{from} = \log(\text{lam0} \times 0.99), \text{to} = \log(\text{lam0} \times 0.1), \text{length} = 10))$$

The runtime comparison among algorithms is reported in Table 1. For imputeFAMD and softImpute, the reported time is the total runtime under all searched hyperparameters.

### 2.2 MAR and MNAR mechanism

We conduct additional experiments under a MAR mechanism and a MNAR mechanism to test the sensitivity of implemented imputation methods.

For MNAR, we use a self-masking mechanism which assigns samples different missing probability by their own values for each variable. Concretely, suppose we want to mask $\alpha$ percentage entries as missing, then for each variable, we assign a high missing probability $(\alpha + 10\%)$ for samples below the first third quantile, a medium missing probability $(\alpha)$ for samples between the first third and the second third quantile, and a low missing probability $(\alpha - 10\%)$ for samples above the second third quantile.

For MAR, we first randomly select $\frac{1}{3}$ of variables as observed. Then for each of the remaining $\frac{2}{3}$ of variables, its samples receive three different missing probability similar to the MNAR mechanism but based a randomly selected observed variable instead of its own values. To have $\alpha$ missing ratio and compensate that only $\frac{2}{3}$ of variables may be masked as missing, we use $\frac{3\alpha}{2}$ as the normal missing probability, $\frac{3\alpha}{2} + 10\%$ as the high missing probability, and $\frac{3\alpha}{2} - 10\%$ as the low missing probability.

### 2.3 Synthetic experiments supplement

Fig 1 of the main paper reports the results for categorical variables with six categories under MCAR of 30% missing ratio. In Fig 2 and Fig 3, we report the results for different number of categories: three and nine. In Fig 4 and Fig 5, we report the results for different missing ratios under MCAR. In Fig 6 and Fig 7, we report the results under different missing mechanism (MAR and MNAR). In general, these experiments show EGC performs well for both categorical and ordered variables in mixed data as reported in Section 3.1 of the main paper.

### 2.4 Real data experiments supplement

All used datasets are accessed from openml.org through the R package OpenML. Table 1 provides an overview of the used datasets. The prepared dataset does not distinguish categorical and ordinal variables. We do distinguish them according to the variable description whenever available. Some
Figure 1: Imputation error on synthetic mixed data under MCAR of \(30\%\) missing. There are 5 continuous variables, 5 ordinal variables and \(1/3/5\) categorical variables with three categories, reported over 10 repetitions (error bars indicate standard deviation).

Figure 2: Imputation error on synthetic mixed data under MCAR of \(30\%\) missing. There are 5 continuous variables, 5 ordinal variables and \(1/3/5\) categorical variables with nine categories, reported over 10 repetitions (error bars indicate standard deviation).

Figure 3: Imputation error on synthetic mixed data under MCAR of \(20\%\) missing. There are 5 continuous variables, 5 ordinal variables and \(1/3/5\) categorical variables with six categories, reported over 10 repetitions (error bars indicate standard deviation).
Figure 4: Imputation error on synthetic mixed data under MCAR of 40% missing. There are 5 continuous variables, 5 ordinal variables and $1/3/5$ categorical variables with six categories, reported over 10 repetitions (error bars indicate standard deviation).

Figure 5: Imputation error on synthetic mixed data under MAR of 30% missing. There are 5 continuous variables, 5 ordinal variables and $1/3/5$ categorical variables with six categories, reported over 10 repetitions (error bars indicate standard deviation).

Figure 6: Imputation error on synthetic mixed data under MNAR of 30% missing. There are 5 continuous variables, 5 ordinal variables and $1/3/5$ categorical variables with six categories, reported over 10 repetitions (error bars indicate standard deviation).
Table 2: Used UCI dataset overview.

<table>
<thead>
<tr>
<th>OpenML ID</th>
<th>n</th>
<th>p_cat</th>
<th>p_ord</th>
</tr>
</thead>
<tbody>
<tr>
<td>Abalone</td>
<td>1557</td>
<td>4177</td>
<td>1</td>
</tr>
<tr>
<td>Heart</td>
<td>53</td>
<td>270</td>
<td>3</td>
</tr>
<tr>
<td>CMC</td>
<td>23</td>
<td>1473</td>
<td>1</td>
</tr>
<tr>
<td>Creditg</td>
<td>31</td>
<td>1000</td>
<td>8</td>
</tr>
<tr>
<td>Credita</td>
<td>29</td>
<td>690</td>
<td>4</td>
</tr>
<tr>
<td>Colic</td>
<td>25</td>
<td>368</td>
<td>4</td>
</tr>
</tbody>
</table>

Figure 7: Imputation error of categorical variables and of ordered variables, i.e., ordinal and continuous, on 6 UCI datasets under MCAR of 10% missingness. Results shown as mean ± standard deviation.

Features are removed because their distribution are highly concentrated at a single value (more than .95%). All preprocessing information are provided in the codes.

Fig 1 of the main paper reports the results under MCAR of 20% missing ratio. In Fig. 7 and Fig. 8 we report the results for different missing ratios under MCAR. In Fig. 9 and Fig. 10 we report the results under different missing mechanism (MAR and MNAR). In general, these experiments show EGC performs well for both categorical and ordered variables in mixed data as reported in Section 3.2 of the main paper.

Figure 8: Imputation error of categorical variables and of ordered variables, i.e., ordinal and continuous, on 6 UCI datasets under MCAR 30% missingness. Results shown as mean ± standard deviation.
Figure 9: Imputation error of categorical variables and of ordered variables, i.e., ordinal and continuous, on 6 UCI datasets **under MAR 20% missingness**. Results shown as mean ± standard deviation.

Figure 10: Imputation error of categorical variables and of ordered variables, i.e., ordinal and continuous, on 6 UCI datasets **under MNAR 20% missingness**. Results shown as mean ± standard deviation.