

# QI Mathematica Package – Brief Manual

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The following is a brief description of a selection of Mathematica commands that may be of use to people studying quantum information. I make no guarantee that they are well written, free of bugs, optimized for speed and memory usage, etc. Most are functions that I have written because I have found them useful, and I have used them and tested them to a reasonable extent (i.e., until I have become confident in them, at least in the instances in which I have employed them). If you find any bugs, I would be pleased to hear of them. As far as I am aware, all commands work in Mathematica 12.

## I. INSTALLATION

No special installation is needed. The file `QI.m` can be downloaded from <https://github.com/rogercolbeck/QI>. Simply open the file and press Run All Code. However, if you wish to automatically load these functions on starting Mathematica:

1. Change to your autoload directory (usually the Autoload sub-directory of your user addons directory, found using the command `$UserAddOnsDirectory`).
2. Create a subdirectory `QI`.
3. Put the file `QI.m` in this directory.
4. Rename the file `init.m`.

A file, [https://www-users.york.ac.uk/~rc973/QI/QI\\_examples.nb](https://www-users.york.ac.uk/~rc973/QI/QI_examples.nb) collects some simple examples illustrating the use of these functions. A palette, <https://www-users.york.ac.uk/~rc973/QI/Palette.nb> can also be downloaded which contains some useful symbols (including  $\otimes$ ).

## II. GENERAL SYNTAX

In order to use these commands, all vectors must be entered as matrices (this makes them of the right “shape” that the same commands (such as tensor product) can be used for both vectors and matrices without modification). That is, to assign the vector  $\begin{pmatrix} 0 \\ 1 \end{pmatrix}$  to `v` one would enter `v = {{0}, {1}}`. It is easy to convert a standard Mathematica vector (i.e., one entered as `v = {0, 1}`, which I call a list) to this form by using `v = Transpose[{v}]`.

The error handling of these commands is basic (although some checks are done, mostly any displayed errors come out of Mathematica’s built-in functions).

Many of the commands use `Chop` for comparisons of numerical values. By default, `Chop` treats numbers smaller than  $10^{-10}$  as 0. This means that some commands may do the wrong thing with numerical inputs that are smaller than  $10^{-10}$ .

## III. COMMAND LIST

### A. Manipulating Quantum States

`CT[M]`: Equivalent to `ConjugateTranspose[M]`.

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`KetV[i, d]`: This creates the  $i^{\text{th}}$  basis vector in dimension  $d$ , where  $i \in \{0, \dots, d - 1\}$ .

`BraV[i, d]`: This creates the  $i^{\text{th}}$  dual basis vector in dimension  $d$ , where  $i \in \{0, \dots, d - 1\}$ . `BraV[i, d]` is equivalent to `CT[KetV[i, d]]`.

`DM[vec]`: Converts the vector `vec` to a density matrix.

`CircleTimes[A, B] ( $\otimes$ )`: This performs the tensor product and can be entered as either  $A \otimes B$  or `CircleTimes[A, B]`. The  $\otimes$  symbol can be found in the operators→general section of the complete characters palette, or may be entered using the keyboard sequence `ESCAPE, c, *, ESCAPE`.

`Tensor[A, B, pos, desc]`: This perform the tensor product of  $A$  and  $B$ , placing the systems in the positions specified by `pos` (a string of 1s and 2s) where the systems have dimensionalities specified by `desc` (note that `desc` does NOT specify the dimensionality of the matrices  $A$  and  $B$ ). E.g. `Tensor[A, B, {1, 2, 1, 2}, {{a1, a2}, {b1, b2}, {c1, c2}, {d1, d2}}]` requires  $A$  to be the tensor product of a  $a_1 \times a_2$  matrix with a  $c_1 \times c_2$  matrix, i.e., to have dimension  $a_1 c_1 \times a_2 c_2$ , and  $B$  to be the tensor product of a  $b_1 \times b_2$  matrix with a  $d_1 \times d_2$  matrix. The output has  $A$  on the 1st and 3rd systems and  $B$  on the 2nd and 4th systems. If `desc` is a simple list, e.g., `{a1, b1, c1, d1}` then  $A$  and  $B$  must be either be square matrices of size  $a_1 c_1$  and  $b_1 d_1$  or vectors of size  $a_1 c_1$  and  $b_1 d_1$ .

`TensorPower[M, n]`: This raises matrix  $M$  to the  $n$ -th tensor power, i.e., computes  $M^{\otimes n}$ .

`ExchangeSystems[A, newpos, desc]`: This takes an operator representing systems in one order and converts it to another, as specified by `newpos`. The first entry of `newpos` gives the position that the first system moves to, etc.  $A$  can be a vector or a matrix. `desc` should be a list of the dimensions, e.g., `desc={{2, 4}, {3, 5}}` means that  $A$  is considered as the tensor product of a  $2 \times 3$  matrix with a  $4 \times 5$  matrix (so  $A$  should have dimensions  $8 \times 15$ ). If the subsystems are square, or are vectors, `desc` need only be the number of rows. For instance, `ExchangeSystems[v1⊗v2⊗v3, {3,1,2}, {2,3,4}]` gives `v2⊗v3⊗v1`, where `v1` has dimension  $2 \times 1$ , `v2` has dimension  $3 \times 1$  and `v3` has dimension  $4 \times 1$ . [Or `v1` has dimension  $2 \times 2$ , `v2` has dimension  $3 \times 3$  and `v3` has dimension  $4 \times 4$ .]

One can also do `ExchangeSystems[DM[v1⊗v2⊗v3], {3,1,2}, {2,3,4}]`, which is the same as `DM[v2⊗v3⊗v1]`.

`DirectSum[A,B,C,...]`: This computes the direct sum of the square matrices  $A, B, C, \dots$ , i.e.  $A \oplus B \oplus C \oplus \dots$

`QubitPartialTrace[M, {i,j,...}]`: This traces out the  $i, j, \dots$  qubits of  $M$ , assuming that  $M$  is the tensor product of multiple qubit spaces.

`PartialTrace[M, dim1, dim2, tr]`: This traces out the  $\text{tr} \in \{1, 2\}$  subsystem of  $M$ , where  $M$  has the Hilbert space structure  $\mathcal{H}_1 \otimes \mathcal{H}_2$ , where  $\mathcal{H}_1$  is  $\text{dim}_1$ -dimensional and  $\mathcal{H}_2$  is  $\text{dim}_2$ -dimensional.

`PT[M, keep, desc]`: This traces out the systems specified by `keep` of  $M$ , where the dimensions of the subsystems in  $M$  are specified by `desc`. The argument `keep` should be a binary vector where 0s are systems to be traced out and 1s are systems to keep. The description `desc` should be a list of the dimensions of the subsystems e.g. `keep={1, 0, 1, 0}`, `desc={2, 4, 2, 3}` traces out the 2nd and 4th systems of  $M$ , where  $M$  is an operator on  $\mathbb{C}^2 \otimes \mathbb{C}^4 \otimes \mathbb{C}^2 \otimes \mathbb{C}^3$  (i.e. a  $48 \times 48$  matrix). The output is a  $4 \times 4$  matrix.

`PTrans[M, action, desc]`: This takes the partial transpose of the systems specified by `action` of  $M$ , where the dimensions of the subsystems in  $M$  are specified by `desc`. The argument `action` should be a binary vector where 1s are systems to be transposed and 0s are systems to do nothing to. The description `desc` should be a list of the dimensions of the subsystems e.g. `action={1, 0, 1, 0}`, `desc={2, 4, 5, 3}` transposes the 1st and 3rd systems of  $M$ , where  $M$  is an operator on  $\mathbb{C}^2 \otimes \mathbb{C}^4 \otimes \mathbb{C}^5 \otimes \mathbb{C}^3$  (i.e. a  $120 \times 120$  matrix).

`BasisForm(S)[vec, desc]`: This writes out a vector, `vec`, with structure given by `desc` in the computational basis, giving one line for each non-zero component (the `BasisFormS` version gives the output all on one line) e.g. `BasisFormS[{1,0,0,1},{2,2}]` returns  $+1|0, 0\rangle + 1|1, 1\rangle$  i.e. it is the state  $|00\rangle + |11\rangle$ , while `BasisFormS[{1,0,0,1},{4}]` returns  $+1|0\rangle + 1|3\rangle$ . Note: this command applies `Flatten` to `vec`, so will do the wrong thing with matrix inputs.

`Purify[rho]`: This gives a state that purifies `rho` (outputting as a vector). The chosen purification takes a copy of the original eigenvector.

**SchmidtDecomposition[v, desc]:** This gives the Schmidt decomposition of the state  $v$  where `desc` is a two component list of the dimensions across which the decomposition is taken. E.g., if  $v$  has eight dimensions, then `desc` can be  $\{2, 4\}$  or  $\{4, 2\}$ . The output is in the form  $\{c, \text{vecs}_1, \text{vecs}_2\}$ , where  $c$  is a list of Schmidt coefficients,  $\text{vecs}_1$  are the Schmidt vectors on system 1, and  $\text{vecs}_2$  are those on system 2 (in the same order as the coefficients in  $c$ ). In other words,  $\text{Sum}[c[[i]] * \text{vecs}_1[[i]] \otimes \text{vecs}_2[[i]], \{i, 1, \text{Min}[\text{desc}]\}]$  is equal to  $v$ .

**DiagonalizingUnitary[M]:** For Normal matrix  $M$ , (i.e.,  $MM^\dagger = M^\dagger M$ ) this returns  $\{U, D\}$ , where  $M = UDU^\dagger$ ,  $U$  is unitary, and  $D$  is diagonal.

**EigensystemExact[A, (prec)]:** For matrix  $A$ , this returns  $\{\text{vals}, \text{vecs}\}$ , where  $\text{vals}$  are the eigenvalues and  $\text{vecs}$  are the corresponding eigenvectors, ensuring that the latter are orthonormal for exact inputs (Mathematica's `Eigensystem` does not do this by default). Note that doing so can be slow. Note also that this command may not behave correctly with symbolic inputs. The eigenvalues are sorted using private function `OrderingF`, hence by the size of their real part, followed by the size of their imaginary part (taking these parts to be equal if within `prec` of each other). If all the eigenvalues are real, they are sorted from smallest to largest, for example. The optional argument `prec` can be used to specify the tolerance for deciding whether eigenvalues are equal (by default this is set to that used by `Chop[]`).

**SimultaneouslyDiagonalize[A, B, (prec)]:** For commuting normal matrices  $A, B$ , this returns  $U$  such that  $U$  is unitary and  $U^\dagger A U$  and  $U^\dagger B U$  are diagonal. This has an optional third argument, which is the precision to which two numerical values are assumed to be equal. By default, this is the standard for `Chop[]`, i.e.,  $10^{-10}$ .

**BlochSphere[ρ]:** This gives the point on the Bloch sphere specified by  $\rho$ , in the form of the vector  $\{r_1, r_2, r_3\}$  satisfying  $\rho = \frac{1}{2} (\mathbb{1} + r_1 \sigma_1 + r_2 \sigma_2 + r_3 \sigma_3)$ .

**FromBlochSphere[r]:** Gives the density matrix corresponding to this point on the Bloch Sphere (see `BlochSphere`).

**AppendCols[U]:** Takes an isometry  $U$  and returns a unitary matrix by completing the columns.

**FillZero[M]:** Takes a matrix with more columns than rows and makes it square by adding rows of zeros.

**MeasureBasis[ρ, basis]:** Performs the quantum channel on  $\rho$  equivalent to measuring in the basis specified by `basis` then forgetting the outcome [to remove the system, use `MeasurePOVM`]. The input `basis` is expressed as a list of vectors, e.g. `basis = {{1, 1}, {1, -1}}/√2` measures in the  $\sigma_x$  basis. Note that this command can be used to measure the  $A$  system of a matrix  $\rho_{AB}$  if the dimensions of  $\rho$  and `basis` do not match. In other words, if the basis vectors are  $\{v_i\}$ , this computes  $\sum_i |v_i\rangle\langle v_i| \rho |v_i\rangle\langle v_i|$  (or  $\sum_i (|v_i\rangle\langle v_i| \otimes \mathbb{1}_B) \rho_{AB} (|v_i\rangle\langle v_i| \otimes \mathbb{1}_B)$ ).

**MeasureBasis[ρ, basis, sys, desc]:** Performs the quantum channel on  $\rho$  equivalent to measuring in the basis specified by `basis` on the `sys` part of  $\rho$  which is partitioned as specified in `desc` then forgetting the outcome [to remove the system, use `MeasurePOVM`]. The input `basis` is expressed as a list of vectors, e.g. `basis = {{1, 1}, {1, -1}}/√2` measures in the  $\sigma_x$  basis. The input `sys` should be given as a vector of 1s and 2s, where 1 is the system to measure. If the third and fourth arguments are blank, the default is to measure the  $A$  system of a matrix  $\rho_{AB}$ , where  $A$  is assumed to have the same dimensions as the specified basis.

**MeasurePOVM[ρ, POVM, sys, desc]:** Performs the quantum channel on  $\rho$  equivalent to measuring the POVM specified by `POVM` on the `sys` part of  $\rho$  which is partitioned as specified in `desc` then tracing out the measured system. The input `POVM` is expressed as a list of matrices (the POVM elements), e.g. `POVM = {{{1, 0}, {0, 0}}, {{0, 0}, {0, 1}}}` measures in the  $\sigma_z$  basis. The input `sys` should be given as a vector consisting of 2s and one 1, where the 1 corresponds to the system to measure. For a bipartite matrix with `sys = {1, 2}` and `desc = {2, 2}`, this computes  $\sum_x |x\rangle\langle x| \otimes \text{tr}_A((E_x \otimes \mathbb{1}) \rho_{AB})$ .

**POVMIsometry[ρ, POVM, sys, desc]:** Performs the isometry on the `sys` part of  $\rho$  equivalent to  $\sum_x |x\rangle\langle x| \otimes |x\rangle \otimes \sqrt{E_x}$ . The input `sys` should be given as a vector consisting of 2s and one 1, where the 1 corresponds to the system to measure. Tracing out the first part of the result together with the part of `sys` corresponding to a 1, is the same as `MeasurePOVM[ρ, POVM, sys, desc]`.

**ChoiState[channel]:** Computes the Choi state of the channel `channel`, which should be specified as a list  $\{K_1, K_2, \dots, K_n\}$  of Kraus operators (all that is needed is that each  $K_i$  is a matrix of the same shape). The output is  $\sum_i (\mathbb{1} \otimes K_i) |\gamma\rangle\langle\gamma| (\mathbb{1} \otimes K_i)$ , where  $|\gamma\rangle = \sum_j |j\rangle \otimes |j\rangle$ .

**ChoiState[set1, set2]:** Computes the Choi state of the linear map defined by `set1` and `set2`, which should be specified as lists  $\{K_1, K_2, \dots, K_n\}$  and  $\{J_1, J_2, \dots, J_n\}$  (all that is needed is that each  $K_i$  and  $J_i$  is a matrix of the same shape and that there are the same number of entries of each). The output is  $\sum_i (\mathbb{1} \otimes K_i) |\gamma\rangle\langle\gamma| (\mathbb{1} \otimes J_i^\dagger)$ , where  $|\gamma\rangle = \sum_j |j\rangle \otimes |j\rangle$ .

**ChoiChannel[state, d<sub>A</sub>, d<sub>B</sub>]:** Computes the Choi channel for the Choi state `state`. The output is  $\{\text{set1}, \text{set2}\}$ , where `set1` and `set2` are lists  $\{K_1, K_2, \dots, K_t\}$  and  $\{J_1, J_2, \dots, J_t\}$ , where  $t$  is at most  $d_A d_B$  and each  $K_i$  and  $J_i$  is a  $d_B \times d_A$  matrix (i.e., the channel will be from  $A$  to  $B$ ), such that the corresponding channel is  $\rho \mapsto \sum_i K_i \rho J_i^\dagger$ . If `state` is positive, the output has  $J_i = K_i$  for all  $i$ .

**ChannelCompress[channel]:** Takes a channel specified by a list of Kraus operators and outputs another representation of the same channel, potentially with fewer Kraus operators. For a channel from dimension  $d_A$  to  $d_B$ , the output has at most  $d_A d_B$  operators.

**ExtremeChannelQ[channel, (tol)]:** Checks whether the channel `channel` is extreme or not. Note that this involves computing the rank, so numerical instability can lead to erroneous results. Optional argument `tol` gives the tolerance for deciding when a singular value is zero.

## B. Random Sampling

For many of these commands, adding **R** to the front restricts to reals, and adding **F** restricts to reals and chooses exact values. The fractional (**F**) version requires an additional argument `prec` which must be an integer (`prec` should not be used for the others). This corresponds to the maximum denominator of the fractions chosen, prior to normalization.

**(R/F)PickRandomPsi[n, (.prec)]:** Gives a random  $n$ -dimensional normalized vector. (**R**) for real version, (**F**) for fractional (real) version. This command samples by picking uniformly distributed random numbers between 0 and 1 for the magnitude, and uniformly distributed random numbers between 0 and  $2\pi$  for the phase and then normalizing the generated list. **F** version samples in a different way.

**(R/F)PickRandomPsi2[n, (.prec)]:** Gives a random  $n$ -dimensional normalized vector. (**R**) for real version, (**F**) for fractional (real) version. This command samples by picking a random number between 0 and 1, and square rooting it for the first probability amplitude. It then picks between 0 and the maximum remaining weight, etc. This hence gives states with more coefficients closer to 0 than the first sampling method. The first coefficients of these states also tend to be the largest. **F** version samples in a different way.

**(R)PickRandomPsiHaar[n]:** Gives a random  $n$ -dimensional normalized vector with Haar distribution. (**R**) takes the absolute value of each component.

**(R/F)PickRandomUnitary[n, (.prec)]:** Gives a random  $n \times n$  unitary matrix. The algorithm works by picking  $n$  linearly independent vectors using `PickRandomPsi` and performing a Gram-Schmidt orthogonalization. (**R**) for real version, (**F**) for fractional (real) version.

**(R)PickRandomUnitaryHaar[n]:** Gives a random  $n \times n$  unitary matrix with Haar distribution. The algorithm works by picking a matrix of complex numbers each with normally distributed real and imaginary parts and performing Gram-Schmidt orthogonalization. (**R**) for real version.

**(R/F)PickRandomIsometry[dim<sub>1</sub>, dim<sub>2</sub>, (.prec)]:** Gives a random isometry from  $\text{dim}_1$  to  $\text{dim}_2$ . (**R**) for real version, (**F**) for fractional (real) version.

**(R/F)PickRandomRho[n, (.prec), (.rank)]:** Gives a random  $n \times n$  density matrix. The algorithm picks a random diagonal version, then applies a random unitary to it. (**R**) for real version (**F**) for fractional (real) version, and optional specification of the rank, `rank` as the final argument.

**(R)PickRandomPOVM[dim, num]:** Outputs a random `num` element POVM with dimension `dim`, i.e., a set of `num` matrices  $E_i$  such that  $\sum_i E_i = \mathbb{1}$ . [Fractional (**F**) version not currently implemented.]

**(R)PickRandomMeasurement[*dim*, *num*]:** Outputs a random *num* element measurement with dimension *dim*, i.e., a set of matrices  $K_i$  such that  $\sum_i K_i^\dagger K_i = \mathbb{1}$ . ( $K_i^\dagger K_i$  are the POVM elements.) [Fractional (F) version not currently implemented.]

**(R/F)PickRandomChannel[*dim*<sub>1</sub>, *dim*<sub>2</sub>, *num* (*,prec*):** Outputs the Kraus form of a random *num* element channel from dimension *dim*<sub>1</sub> to dimension *dim*<sub>2</sub>, i.e., a set of *num* matrices  $K_i$  such that  $\sum_i K_i^\dagger K_i = \mathbb{1}$ . In other words, the intended channel is  $\rho \mapsto \sum_i K_i \rho K_i^\dagger$ .

### C. Distance Measures And Distinguishing States

**Dist[ $\rho_1$ ,  $\rho_2$ ]:** This computes the trace distance between matrices  $\rho_1$  and  $\rho_2$ , i.e.  $\frac{1}{2} \text{tr} |\rho_1 - \rho_2|$ .

**QuickDist[ $\rho_1$ ,  $\rho_2$ ]:** This computes the trace distance in a different (usually faster) way. The difference between these two methods is that the second finds the eigenvalues and works on these, while the first relies on Mathematica's built in `MatrixPower` command.

**Fidelity[ $\rho_1$ ,  $\rho_2$ ]:** This computes the fidelity between the two states, i.e.  $\text{tr} \sqrt{\sqrt{\rho_1} \rho_2 \sqrt{\rho_1}}$ .

**OptimumPOVM[ $\rho_1$ ,  $\rho_2$ ,  $\mathbf{p}_1$ ]:** This returns  $\{E_0, E_1\}$ , where these are the POVM elements that optimally distinguish  $\rho_1$  and  $\rho_2$ , where  $\rho_1$  occurs with probability  $\mathbf{p}_1$ , and  $\rho_2$  with probability  $1 - \mathbf{p}_1$ . One can enter `OptimumPOVM[ $\rho_1$ ,  $\rho_2$ ]`, in which case it is assumed that  $\mathbf{p}_1 = \frac{1}{2}$ . The POVM is optimal in the sense that it maximizes the probability of correctly guessing whether a given state is  $\rho_1$  or  $\rho_2$ .

### D. Entropy Measures

The entropy commands in this section may not work for symbolic matrices because they use `If` statements to distinguish cases and the conditions may not be resolvable for symbolic values.

**Matrixlog[M]:** Returns the natural logarithm of matrix *M*, where *M* is Hermitian.

**Matrixlog[b, M]:** Returns the logarithm of matrix *M* in base *b*, where *M* is Hermitian.

**Matrixxlogx[M]:** Returns the function  $x \log x$  applied to Hermitian matrix *M*, taking  $0 \log 0$  to be 0.

**Matrixxlogx[b, M]:** Returns the function  $x \log_b x$  applied to Hermitian matrix *M*, taking  $0 \log_b 0$  to be 0.

**ShanEntropy[*problist*]:** Computes the Shannon entropy of the distribution given by *problist* =  $\{p_1, p_2, \dots\}$ , i.e.  $\sum_i -p_i \log_2 p_i$ .

**ShanEntropy[*p*]:** This is equivalent to `ShanEntropy[{p, 1-p}]`.

**vNEntropy[*rho*, *keep*, *desc*]:** Computes the (conditional) von Neumann entropy of *rho*, a multi-partite system whose dimensions are given by *desc*, conditioning on the systems specified by *keep* which uses 1 for the system, 2 for systems to condition on and 0 for systems to trace out. Example: `vNEntropy[rho, {1,0,2}, {2,4,2}]` takes a  $2 \times 4 \times 2$  dimensional matrix *rho*, traces out the 4 dimensional subsystem, then calculates the von Neuman entropy of the first system given the third.

**vNInfo[*rho*, *keep*, *desc*]:** Computes the (conditional) von Neumann mutual information of *rho*, a multi-partite system whose dimensions are given by *desc*, conditioning on the systems specified by *keep* which uses 1 for the first system, 2 for the second system, 3 for systems to condition on and 0 for systems to trace out. Example: `vNInfo[rho, {1,2,0,3}, {2,4,2,2}]` takes a  $2 \times 4 \times 2 \times 2$  dimensional matrix *rho* and computes the quantum mutual information between the first and second systems given the fourth.

**RelEnt[A, B]:** Computes the von Neumann relative entropy of *A* and *B*, equal to  $(\text{tr}(A \log A) - \text{tr}(A \log B))/\text{tr}(A)$ . If *A* is not contained in the support of *B* then the answer is infinite.

**RelEnt**[ $\alpha$ , A, B]: Computes the  $\alpha$  relative entropy of A and B, equal to  $\frac{1}{\alpha-1} \log_2 \text{tr}(A^\alpha B^{1-\alpha})$ . For  $\alpha = 0$  this evaluates the relative entropy and for  $\alpha = 1$ , the expression is interpreted via the limit  $\alpha \rightarrow 0^+$ .

**RenyiEnt**[ $\alpha$ , rho, keep, desc]: Computes the  $\alpha$  Rényi entropy of rho, a multi-partite system whose dimensions are given by desc, conditioning on the systems specified by keep which uses 1 for the system, 2 for systems to condition on and 0 for systems to trace out. Here the conditional Rényi entropy is defined by  $H_\alpha(A|B) = \frac{1}{1-\alpha} \log_2 \text{tr}(\rho_{AB}^\alpha (\mathbb{1} \otimes \rho_B)^{1-\alpha})$ . For  $\alpha = 1$  this evaluates the von Neumann entropy, and for  $\alpha = 0$ , the expression is interpreted via the limit  $\alpha \rightarrow 0^+$ .

## E. Linear Programming

**(N)RemoveIneqConstraints**[M, b]: Removes redundant inequality constraints from the set of equations specified by matrix M where  $Mx \geq b$ , returning  $\{M', b'\}$ . The parameter b can also be given in the form of pairs of values  $\{\{b_1, s_1\}, \{b_2, s_2\} \dots\}$  where  $s_i \in \{-1, 0, 1\}$  represents less than or equal to, equal to, or greater than (the syntax is the same as in Mathematica's **LinearProgramming** command). Equality constraints are retained. This works by optimizing each constraint subject to the others to see whether it is already implied. (N) is for the numerical version, where each **LinearProgramming** minimization is done numerically (which is faster). It may also be desirable to use **RemoveDuplicateConstraints**[M, b] first to speed up the process.

**RemoveDuplicateConstraints**[M, b]: Removes redundant constraints from the set of equations specified by matrix M where  $Mx \geq b$ , returning  $\{M', b'\}$ . The parameter b can also be given in the form of pairs of values  $\{\{b_1, s_1\}, \{b_2, s_2\} \dots\}$  where  $s_i \in \{-1, 0, 1\}$  represents less than or equal to, equal to, or greater than (the syntax is the same as in Mathematica's **LinearProgramming** command). This works by checking for identical rows in M and processing them accordingly (if  $Mx = 5$ , any inequality constraints  $Mx \geq c$  are removed, if  $Mx \geq 4$  and  $Mx \geq 5$ , then the former is removed). Note that no consistency check is performed on the equations, so an inconsistent set may become consistent, since if  $Mx = 3$  and  $Mx = 4$ , then one of these will be removed.

**Prep**[M, b]: Here b must have the form  $\{\{b_1, s_1\}, \{b_2, s_2\} \dots\}$  (the syntax is the same as in Mathematica's **LinearProgramming** command) and is converted to  $\{M', b'\}$ , where  $M'x \geq b'$ . Thus, if  $s_1$  is negative the sign is flipped and if  $s_1 = 0$ , two inequalities are included corresponding to  $M_i x \geq b_i$  and  $-M_i x \geq -b_i$ .

**FourierMotzkin**[M, b, elim, (Options)]: Here b must have the form  $\{b_1, b_2, \dots\}$  and signifies  $Mx \geq b$  component-wise. The command produces a new set of inequalities eliminating the variables specified by elim, e.g., if  $\text{elim}=\{1,4\}$ , then the first and fourth variables are eliminated. The command returns  $\{M', b'\}$  such that  $M'x' \geq b'$ , where  $x'$  now does not involve the eliminated variables. This command tries to optimize the order of the eliminations (choosing the next variable to keep the total number of inequalities smallest), and uses **RemoveIneqConstraints**[M, b] after each step to reduce the number before starting the next elimination [to turn this off, use the option **Simp** $\rightarrow$ **False**]. To go through the eliminations one at a time without removing redundant equations, use **Elim**[M, b, i] to eliminate the  $i$ th variable only. Note that the **Elim** function adds in a row to the matrix corresponding to the positivity of the element to be removed (such a constraint is assumed by **LinearProgramming**) [to turn this off, use the option **AddId** $\rightarrow$ **False**].

[Commands **NegInstances**[M, i] and **PosInstances**[M, i] are private for **Elim**.]

## F. Miscellaneous

**IntDigs**[num, bases]: Writes the number num in terms of the bases specified in bases. The length of the output is always equal to the number of bases. E.g., **IntDigs**[16, {4,2,3}] gives {2, 1, 1} meaning that  $16 = 2 \times (2 \times 3) + 1 \times 3 + 1 \times 1$ . Note that **IntDigs**[24, {4,2,3}] gives {0, 0, 0} since the most significant digit cannot be larger than 4. The next digit would give the number of  $4.3.2=24s$ .

**Progress**[i, i\_min, i\_max, num]: This command is useful to give a progress report in large runs. Assuming i increases by an integer each time, and runs between i\_min and i\_max, this will display num outputs throughout this range. For example, to get a progress indication every 10%, use **num**=10. [N.B. This command may also work if i increments by more than an integer each time, provided num is not too large.]

`ProgressTemporary[i, i_min, i_max, num]`: As above but uses `PrintTemporary` so that the progress display disappears on completion.

`(N)ThreadSolve[eqns, soln, var]`: Solves the set of equations `eqns[[i]] = soln` for variables `var`. Example: `ThreadSolve[{a + 1/2, 6 * a + 3/2}, 0, a]` returns `{-1/2, -1/4}`, these being the values of `a` that solve the first and second equations. `(N)` is for the numerical version.

`$\sigma$ [i]`: This outputs the  $i^{\text{th}}$  Pauli Matrix, with  $i \in \{0, 1, 2, 3\}$ . The  $\sigma$  symbol is in the basic calculations palette, or can be entered using the keyboard sequence `ESCAPE, s, ESCAPE`.

`Support[M]`: Returns the projector onto the support of a matrix `M`.

`PosPart[M]`: Returns the (strictly) positive part of a matrix `M`. Do not use with symbolic matrices.

`PosPart[M, assum]`: Returns the (strictly) positive part of a matrix `M`, making assumptions `assum` about certain variables in `M`. Example: `PosPart[{{Sin[t]^2, -Sin[t]*Cos[t]}, {-Sin[t]*Cos[t], -Sin[t]^2}}`] returns `{0, 0}` with a warning message, but `PosPart[{{Sin[t]^2, -Sin[t]*Cos[t]}, {-Sin[t]*Cos[t], -Sin[t]^2}}, 0 | t | 1]` returns a meaningful answer. Care is needed for use with symbolic matrices.

`NegPart[M]`: Returns the (strictly) negative part of a matrix `M`. Do not use with symbolic matrices.

`NegPart[M, assum]`: Returns the (strictly) negative part of a matrix `M`, making assumptions `assum` about certain variables in `M`. Care is needed for use with symbolic matrices.

`AntiDiagonalMatrix[M]`: Turns a matrix into a vector of diagonal values. Example: `AntiDiagonalMatrix[{{a, b}, {c, d}}]` returns `{{a}, {d}}`. Note: Mathematica's `Diagonal` command is similar, up to the form of the output.

`CreateMatrix[W, r, c]`: This creates the  $r \times c$  matrix with elements `W[i, j]`.

`CreateHermitianMatrix[W, r, c]`: This acts like `CreateMatrix` except that it ensures the output is Hermitian. May be appended to `CreateHermitianMatrixR[W, r, c]` to further ensure that the variables used are real, i.e. the elements are now `WR[i, j] + i WC[i, j]` (and `WC[i, i] = 0`).

`CreateSymmetricMatrix[W, r, c]`: As `CreateHermitianMatrix[W, r, c]` but the matrix created is symmetric rather than Hermitian.

`RemoveRepeatedInstances[list]`: This returns a smaller list which features only one instance of each entry. The order is preserved based on the order in which the first instance occurred in the original list. Example: `RemoveRepeatedInstances[{a, b, b, a, 1, s, 1}]` returns `{a, b, 1, s}`. Note: Mathematica's `DeleteDuplicates` command also does this.

`SetMinus[S1, S2]`: If `S1` and `S2` are two lists, this computes `S1 \ S2`, i.e. it returns `S1` with the elements of `S2` removed. Note that if `S1` is not a set, e.g., it contains repeated entries, then so does the output. The order is maintained. Example: `SetMinus[{1, 2, 3, 4}, {2, 3}]` returns `{1, 4}`.

`MajorizeQ[A, B]`: Checks whether `A` majorizes `B`, where `A` and `B` are lists of numbers, i.e. whether when ordered by size with  $a_1 \geq a_2 \geq \dots$ , and  $b_1 \geq b_2 \geq \dots$ , we have  $a_1 \geq b_1$ ,  $a_1 + a_2 \geq b_1 + b_2$  etc.

`ManualDeriv[fn, vars, point, steps]`: Computes an estimate of the derivative of the function `fn`, which should be a function of the variables in set `vars` at the point `point` with set of stepsizes `steps`. This is useful for complicated expressions that cannot be differentiated symbolically. The function is best input within `Hold[ ]`. E.g., `ManualDeriv[Hold[Integrate[y*x, {x, 0, a}, {y, b, c}]], {a, b, c}, {3, 2, 3}, {1/100, 1/10, 1/10}]` treats the integral as a function of `a, b, c` and uses steps `1/100, 1/10, 1/10`. The first part of the output is `(f[3+1/100, 2, 3] - f[3, 2, 3]) / (1/100)` etc., where `f[a,b,c]` is the integral.

`QRDecomp[M]`: For a square matrix `M`, returns square matrices `{Q, R}` such that `Q` is unitary, `R` is right (upper) triangular and `Q†R = M`. Note: Mathematica's built in `QRDecomposition` does not output square matrices if `M` doesn't have full rank.

**RQDecomp[M]:** For a square matrix  $M$ , returns square matrices  $\{R, Q\}$  such that  $R$  is right (upper) triangular,  $Q$  is unitary and  $RQ^\dagger=M$ .

**QLDecomp[M]:** For a square matrix  $M$ , returns square matrices  $\{Q, L\}$  such that  $Q$  is unitary,  $L$  is left (lower) triangular and  $Q^\dagger L=M$ .

**LQDecomp[M]:** For a square matrix  $M$ , returns square matrices  $\{L, Q\}$  such that  $L$  is left (lower) triangular,  $Q$  is unitary and  $LQ^\dagger=M$ .

**QRDecompPos[M]:** As **QRDecomp** but ensures that  $R$  has positive diagonal entries. [Similarly, **RQDecompPos**, **LQDecompPos** and **QLDecompPos**.]

[These use the private function **SignZeroPos[v]** which takes a list  $v$  and outputs a list of the signs of (the real parts of)  $v$  but taking 0 to have positive sign, e.g. **SignZeroPos**[[1,0,-1,-2+I,0]] gives {1,-1,-1,-1,-1}.]

**QRDecompNeg[M]:** As **QRDecomp** but ensures that  $R$  has negative diagonal entries. [Similarly, **RQDecompNeg**, **LQDecompNeg** and **QLDecompNeg**.]

[These use the private function **SignZeroNeg[v]** which takes a list  $v$  and outputs a list of the signs of (the real parts of)  $v$  but taking 0 to have negative sign, e.g. **SignZeroPos**[[1,0,-1,-2+I,0]] gives {1,-1,-1,-1,-1}.]