An Exploration into Linear Algebra

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It's Too Early for This

1 Basic Matrix Code

To begin, we have will create code that represents matrices and performs elementary matrix operations, such as computing the inverse, multiplying, and adding.

We will utilise Python to do so. Python has a data structure known as a "list" or an "array." These are essentially a collection of indexed data that can be manipulated. Lists may contain sublists; it is in this way that we can represent a "matrix" in python, for indeed, a matrix is nothing but a collection of row/column vectors, which themselves can be represented as an individual python list.

We have decided to represent a matrix as a collection of row vectors. For example consider the following matrix, $A \in \mathbb{R}^{m \times n}$:

$$A = \begin{bmatrix} a_1 & a_2 & a_3 & \dots & a_n \\ b_1 & b_2 & b_3 & \dots & b_n \\ \dots & \dots & \dots & \dots & \dots \\ m_1 & m_2 & m_3 & \dots & m_n \end{bmatrix}$$

with m rows and n columns. We choose to represent the same matrix, A pythonically in the following way:

```
array = [
    [a1, a2, a3, ..., aN],
    [b1, b2, b3, ..., bN],
    [c1, c2, c3, ..., cN],
    ...,
    [m, m2, m3, ..., mN],
]
```

In this way, each sublist is a row of the matrix.

In the following we, will discuss how to code basic matrix operations using this data structure.

1.1 Multiplying a Matrix by a Scalar

One of the most crucial elements of matrix arithmetic is the ability to multiply a given matrix by a scalar. When doing this, the following will hold true:

$$\forall \lambda \in \mathbb{R}, \forall A \in \mathbb{R}^{m \times n}, \lambda A = \begin{bmatrix} \lambda a_1 & \lambda a_2 & \lambda a_3 & \dots & \lambda a_n \\ \lambda b_1 & \lambda b_2 & \lambda b_3 & \dots & \lambda b_n \\ \dots & \dots & \dots & \dots \\ \lambda m_1 & \lambda m_2 & \lambda m_3 & \dots & \lambda m_n \end{bmatrix},$$

This is a rather simple problem to tackle. Simply, we iterate through each row of the matrix, and then through each element (column) within that row, multiplying each entry by a given λ as we iterate. This produces a matrix that has been multiplied by lambda. The code is as follows:

```
def matrix_by_scalar(matrix1, scalar_quantity):
   , , ,
   TAKES:
       A matrix of the form outlined above, matrix1
       A scalar_quantity by which the matrix will be multiplied
   RETURNS:
       A matrix of the form outlined above
   try:
       if (isinstance(scalar_quantity, int)) or
          (isinstance(scalar_quantity, float)):
       # O(1) we wcheck if the scalar element is a valid real
          number; if not, we raise an error.
           return list([element * scalar_quantity for element in
              row] \
              for row in matrix1) # O(n**2) This comprehension
                  multiplies each element of each row by the
                  scalar and thus has a time complexity of n**2
```

```
else: raise ValueError(f"Argument passed:
    '{scalar_quantity}'. Error: Expected argument of type
    'int' or 'float' ") # error raised
except: # In Case something goes wrong; biggest error here is
    the potential for incorrect args passed
    print("something went wrong - likely the matrix argument
        was incorrect")
```

1.2 Adding Two Matrices

The addition of matrices is an elementwise one, and therefore it is easy to implement iteratively and makes lots of intuitaive sense. What is meant by elementwise is that we can handle each element individually; in the case of matrix addition, we add each element of a given matrix to the corresponding element of a separate matrix. We can assert that the matrices must therefore be the of the same dimension to be added.

Essentially, we have the following for $\forall A, B \in \mathbb{R}^{m \times n}$, where i = 1, ..., m and j = 1, ..., n

$$A+B = \begin{bmatrix} A_{1,1} & A_{1,2} & \dots & A_{1,j} \\ A_{2,1} & A_{2,2} & \dots & A_{2,j} \\ \dots & \dots & \dots & \dots \\ A_{i,1} & A_{i,2} & \dots & A_{i,j} \end{bmatrix} + \begin{bmatrix} B_{1,1} & B_{1,2} & \dots & B_{1,j} \\ B_{2,1} & B_{2,2} & \dots & B_{2,j} \\ \dots & \dots & \dots & \dots \\ B_{i,1} & B_{i,2} & \dots & B_{i,j} \end{bmatrix}$$

$$= \begin{bmatrix} A_{1,1} + B_{1,1} & A_{1,2} + B_{1,2} & \dots & A_{1,j} + B_{1,j} \\ A_{2,1} + B_{2,1} & A_{2,2} + B_{2,2} & \dots & A_{2,j} + B_{2,j} \\ \dots & \dots & \dots & \dots \\ A_{i,1} + B_{i,1} & A_{i,2} + B_{i,2} & \dots & A_{i,j} + B_{i,j} \end{bmatrix}$$

The code looks as follows:

```
if (len(mat1) == (len(mat2)) and (len(mat1[0]) ==
   len(mat2[0]))): # check to see if possible to add the two
   lists, as their dimensions must be the same
   The following is just a long list comprehension that
       iterates through each matrix and adds corresponding
       elements. It then appends these to "l", which is what is
       ultimately what is returned
   , , ,
   # basically just two for loops
   1 = list([mat1[row][col]+mat2[row][col] \
       for col in range(len(mat1[0]))] \
       for row in range(len(mat1)))
   return 1
   # Intuitively subtraction is very similar: the "+" must be
       turned to a "-"
else: # if the matrices are not of the same size.
   raise ValueError("Args are different size and thus cannot
       be added")
```

This code will hold for subtraction as well, because of the following identity:

$$A - B = \begin{bmatrix} A_{1,1} & A_{1,2} & \dots & A_{1,j} \\ A_{2,1} & A_{2,2} & \dots & A_{2,j} \\ \dots & \dots & \dots & \dots \\ A_{i,1} & A_{i,2} & \dots & A_{i,j} \end{bmatrix} - \begin{bmatrix} B_{1,1} & B_{1,2} & \dots & B_{1,j} \\ B_{2,1} & B_{2,2} & \dots & B_{2,j} \\ \dots & \dots & \dots & \dots \\ B_{i,1} & B_{i,2} & \dots & B_{i,j} \end{bmatrix}$$

$$= \begin{bmatrix} A_{1,1} - B_{1,1} & A_{1,2} - B_{1,2} & \dots & A_{1,j} - B_{1,j} \\ A_{2,1} - B_{2,1} & A_{2,2} - B_{2,2} & \dots & A_{2,j} - B_{2,j} \\ \dots & \dots & \dots & \dots \\ A_{i,1} - B_{i,1} & A_{i,2} - B_{i,2} & \dots & A_{i,j} - B_{i,j} \end{bmatrix}$$

where $\forall A, B \in \mathbb{R}^{m \times n}$, and i = 1, ..., m, j = 1, ..., n.

This implies that we must merely change the elementwise addition in our code to subtraction in order to create a function that subracts two matrices.

1.3 Transpose of a Matrix

The transpose of a matrix is a modification to its structure such that rows become columns and vice versa. That is, the first row of a given matrix A will be the first column of it's transpose, A^T , the second row of A will be the second column of A^T , and so forth and so on.

$$A = \begin{bmatrix} A_{1,1} & A_{1,2} & \dots & A_{1,j} \\ A_{2,1} & A_{2,2} & \dots & A_{2,j} \\ \dots & \dots & \dots & \dots \\ A_{i,1} & A_{i,2} & \dots & A_{i,j} \end{bmatrix} \in \mathbb{R}^{m \times n}, i = 1, \dots, \text{ and } j = 1, \dots, n$$

Then,

$$A^{T} = \begin{bmatrix} A_{1,1} & A_{2,1} & \dots & A_{i,1} \\ A_{1,2} & A_{2,2} & \dots & A_{i,2} \\ \dots & \dots & \dots & \dots \\ A_{1,j} & A_{2,j} & \dots & A_{i,j} \end{bmatrix} \in \mathbb{R}^{m \times n}, i = 1, \dots, \text{ and } j = 1, \dots, n$$

In the case of square matrices, we can state the matrix has been "flipped" over it's main diagonal, or the diagonal elements running from top-left to bottom-right.

The transpose, as with matrix addition/subtraction and multiplying by a scalar, is rather trivial to implement. We iterate through the columns of a matrix, and return them to the user as rows of a new matrix, which is the transpose. In regards to greabbing the columns of a matrix: python provides no way to index a column as simply as one can index a row. This is because, as we have represented matrices with row vectors, the columns are comprised of elements from multiple different rows. Therefore, we construct a function that enables us to quickly "grab" a column from a matrix given its index.

```
def get_col(matrix_2d, _index):
    return list(row[_index] for row in matrix_2d) # O(n) this
        simply grabs the column from the specified index.

# full time O(n)

def transpose(matrix):

new_array = [get_col(matrix, i) for i in range(len(matrix[0]))]
    # O(n) and nested O(n), becomes O(n**2). takes a column and
    makes it a row
```

1.4 Row Reduction: Echelon/Upper Triangle Form

One of the most important matrix-manipulations is the ability to perform elementary "row operations" upon a matrix. That is, multiplying a row by a scalar, swapping the position of two rows in a matrix, and adding/subtracting a scalar multiple of one row to/from another. These operations have incredibly useful properties, and are widely used to calculate determinants and inverses of matrices, and additionally are used to solved systems of linear equations using matrices.

These row operations allow one to effectively mutate a matrix into one of many various forms; for example, we can row reduce such that what we are left with is of the "upper triangle" form, in which all elements not above or in the main diagonal are 0.

This form is known as "echelon" form, and the process to find it is called "Gaussian Elimination."

We begin the algorithm by swapping rows that need to be swapped.

```
if all((i == 0) for i in col[col_index:]): #0(n) if the
   entire column is filled with zeroes, we call
   continue and the program returns to the initial for
   loop, and goes to the next column
   continue
elif col[col_index] == 0: # 0(1) if one of the elements
   on the diagonal is zero - this is where the dividing
   by zero error occurs so we need to handle this
   , , ,
   here we iterate through all of the rows below the
      diagonal.
   if we find a row that doesn't contain a zero in the
      diagonal column index,
   we will swap them
   , , ,
   for i in range(len(col[col_index:])): #O(n)
       if col[col_index:][i] != 0: # 0(1)
          row_idx = col_index+i # 0(1)
          break
   # the below line of code simple swaps the rows
   matrix[col_index], matrix[row_idx] =
      matrix[row_idx], matrix[col_index] # 0(1)
```

Now that rows have been sufficiently swapped, we may begin subtracting scalar multiples of rows from rows below it. This enables us to achieve a zero in the desired spot. We will continue in the for loop that we currenly operate in.

```
3
                             0
     7
                             0
     2
                             0
       ]
                               ]
we can achieve this via subtracting a scalar multiple of
   the row such that we get 0
ex: row 2. we can achieve row 1, col 2 equaling 0 by
subtracting row 1 *
   matrix[row1_idx] [col2_idx]/matrix[row1_idx] [row_idx]
for row_index in range(len(col)): # O(n) we iterate through
   row of each colum we have grabbed earlier
   ,,,
   remember, we only want to turn the rows below
   the diagonal into 0. thus, we check if the row is indeed
       one we one to turn into 0
   if it is not, its idx will be less than the column idx
   if that proves to be true, we will simple pass
   if row_index <= col_index: #0(1)n checks if the row is
       one we do not want to turn to 0
       ,,,
       the following if statement is unnecessary, as I
          could have explicity called:
          matrix[col_index][col_index] when I called
              denominator later
          matrix[col_index] when I call
              raw_subtractant_row later
          however, will keep this code for readability, as
              I find this easier to understand.
       , , ,
       if row_index == col_index: #0(1)
          denominator = matrix[row_index][col_index] #0(1)
          raw_subtractant_row = matrix[row_index] #0(1)
       pass
   else:
       here we actually do the conversion to 0
```

4

return matrix

1.5 Row Reduction: Row Echelon Form (REF)

Consider a matrix that has been reduced such that it is in echelon form, where all elements below the main diagonal are 0. It is possible to reduce a matrix further. Specifically, by identifying all non-zero diagonal elements and dividing the rows in which they appear by the element itself, we can get to a form similar to echelon form, but one where all non-zero diagonal elements will be 1. This is "row-echelon" form.

In other words, we can obtain the following, given $A \in \mathbb{R}^{m \times n}$ and it is in echelon form:

$$A = \begin{bmatrix} A_{1,1} & A_{1,2} & \dots & A_{1,j} \\ 0 & A_{2,2} & \dots & A_{2,j} \\ \dots & \dots & \dots & \dots \\ 0 & 0 & \dots & A_{i,j} \end{bmatrix}$$

$$A = \begin{bmatrix} 1 & A_{1,2} & \dots & A_{1,j} \\ 0 & 1 & \dots & A_{2,j} \\ \dots & \dots & \dots & \dots \\ 0 & 0 & \dots & 1 \end{bmatrix}$$

It it is easy to turn a matrix in echelon form into one in row-echelon, as we must simply divide by the reciprocal of the diagonal element.

1.6 Row-Reduced Echelon Form

1.7 Finding Pivot Columns

As we discussed earlier in the RREF (Reduced Row Echelon Form) section, our result from that operation is a matrix taken from upper triangle form to one that has ones along its diagonal. In some cases, the matrix has linearly dependent columns, in that case, we want to find the index of those columns for all types of matrices, no matter the size. We iterate through each row to find the first appearance of a 1 and check all prior elements to make sure they are 0's to account for non-square matrices.

```
def identify_pivots(mat, rounded=True):
# takes matrix of form outlined in flowerbox
# returns a list of all indexes in which there is a pivot
   column upon row reducing the input matrix
# identifies the pivot columns of the matrix by row reducing
matrix = mat.copy() # we make a copy because otherwise the
   original matrix is modified and we get fucked
matrix = ref(matrix) # we run ref to row reduce it. we do not
   care about full row reduction because that it is trivial in
   this case
if rounded: # this combats error where 0.99999999999 is not
   read as one, and therefore not read as a pivot
   matrix = mround(matrix, 12)
pivot_col_idx_list = [] # we initialize a blank list will
   contain indexes of pivot cols
for row in matrix: # we iterate through each row
   try: # we add a try in case there are no ones in the row
       first_one = row.index(1) # we find the first appearance
          of a 1 within the list
       for idx2 in range(first_one): # we then check if all
          prior elelements are 0
           if row[idx2] != 0: # if one is not a 0, we contiknue
              the loops
              continue
```

```
# however, if all elements prior to the first 1 are 0,
    we will append the index of that 1 to the pivcollist
pivot_col_idx_list.append(first_one)
```

except: # if there are no ones, it can be infered that
 there is no pivot column in that row, so we continue
 continue

return pivot_col_idx_list

O(n**2) time

1.8 Finding the Determinant

1.9 Inverting a Matrix

1.10 Multiplying Two Matrices

2 Vector(Sub)Spaces Pythonically

2.1 General Introduction to Vector Subspaces

A vector space is special type of a group that contains both an inner and outer operation, which are addition and multiplication by a scalar, respectively. Subspaces are, by proxy, spaces that retain the same properties and are a part of the original vector space

Early on, we realized that it was impossible to represent an infinite series of elements as a vector space in python, for example \mathbb{R}^2 . We realized that we would have to have the user to insert a matrix representing that vector space. We created an umbrella class called "subspace". These are integral to representing linear mappings pythonically.

Our subspace class is very broad, the operations done in the class are more of a general operations as stated below.

2.2 Issues when Representing Subspaces Pythonically

As I said earlier, we cannot express common vector spaces used in linear algebra and we had to settle for finite spaces. They would be in the form of:

$$V = \begin{bmatrix} A_{1,1} & A_{1,2} & \dots & A_{1,j} \\ A_{2,1} & A_{2,2} & \dots & A_{2,j} \\ \dots & \dots & \dots & \dots \\ A_{j,1} & A_{j,2} & \dots & A_{j,j} \end{bmatrix} \in \mathbb{R}^{j \times j}$$

j represents the general dimensionality of the vector space.

2.3 Defining a Space with a Basis

The definition of a basis is the set of linearly independent vectors that span a space. These have many useful applications when it comes to linear mapping.

$$U = span(\begin{bmatrix} 0 \\ 1 \\ 2 \end{bmatrix}, \begin{bmatrix} 1 \\ 1 \\ 2 \end{bmatrix}, \begin{bmatrix} 2 \\ 2 \\ 4 \end{bmatrix})$$

Basis Vector

$$\begin{bmatrix} 0 \\ 1 \\ 2 \end{bmatrix}$$

The code to do this involved taking the indexes of the pivot columns from earlier and then mapping those to the original matrix. In the example above, the pivot columns would be a list with index [0] since the original matrix's RREF form comes out to be

$$\begin{bmatrix}
1 & 0 & 0 \\
0 & 1 & 2 \\
0 & 0 & 0
\end{bmatrix}$$
(1)

def create_basis(self):

we find the basis of generatign set

tempgset = self.gset # we create a temporary gset to not
 modify the original

if self.columned == False: # if the input generating
 vectors are row vecs

Then any valid subspace is just the span of the linearly independent basis vectors.

2.4 Orthonormality of Basis Vectors

Orthonormality of basis vectors is when basis vectors are the unit vectors (the distance of the vector is 1) and if it is orthogonal (purpendicular). We can find this by taking the dot product, a type of inner product defined as $\begin{bmatrix} x & y \end{bmatrix} \cdot \begin{bmatrix} a & b \end{bmatrix} = xa + yb$. The following needs to be satisfied:

```
\langle x , y \rangle = 0
\langle x , x \rangle = 1
```

Here is our dot product code:

```
def dot(v1, v2):
# find the dot product of 2 vectors
if not(isinstance(v1[0], list)) and not(isinstance(v2[0],
    list)) and (len(v1)==len(v2)):
# if we find that the vectors are not 1 dimensional or that
        that they are different sizes
total = 0
    for idx in range(len(v1)):
# multiply elementwise
        total += v1[idx]*v2[idx]
return total
```

These are very important to rotations of vectors. Here is our implemenentation of orthonormality:

```
def isorthonormal(self):
    if self.columned == False: # if the input basis vecs are
```

```
row vecs, a
   basis = mp.transpose(self.basis) # we will make them col
       vecs, but will use a new var to not modify self.basis
else:
   basis = self.basis # we don't want to fuck with
       self.basis
if len(basis) >= 2: # orthonormal must have at least two
   basis vectors
   for idx, vec1 in enumerate(basis): # now we run through
       each basis vector O(n)
       for vec2 in basis[idx+1:]: # we run through each
           other basis vecor
           if (dot(vec1, vec2) != 0) or (dot(vec1, vec1) !=
               1): # if any of them do not pass the test
               (\langle v1, v2 \rangle = 0 \text{ or } \langle v1, v1 \rangle = 1) \text{ (dot defined)}
               lower)
               return False # then we will return False
                   because all must pass the test
   return True # otherwise, if False not returned yet, we
       know it to be true
else: return False # if only one basis vec, nothing to be
   orthonormal to
```

3 Representing Linear Mappings Pythonically

As stated earlier, we would use the user inputted matrix representations of a vector space (generating set) and get a collection of basis vectors. A linear mapping is a function that maps between 2 vector spaces and also prserves the following rules:

```
T: x \to y T(x+y) = T(x) + T(y) T(ax) = aT(x)
```

where T is a function linearly mapping vector space x to y. In the case of a code application, these would be finite vector spaces. To map a vector

from one space to another, we need to find the transition matrix from the basis vectors we are converting with, which could even be 2 bases for each vector space. We will cover that code in the next section. Assuming we have the transition matrix, you can multiply that to the vector to recieve a vector that has been mapped to a new vector space. We can map a given vector with this code:

```
def map(self, vector):
     return self.apply(vector)
def apply_mapping(self, vector):
 # expects a vector in the domain, not codomain
 # expects a vector in the domain, not codomain, of the form
     vector = [element1, ..., element 2]
 # will then apply the linear mapping (self.mapping) to the
 # will return the result
 if len(vector) != self.domain_dim:
   return "hold up! wait a minute! sumn aint right"
 else:
   return mp.multiply_matrix(self.mapping,
       mp.transpose([vector]))
def is_subspace(self, subspace):
   # check if subspace is a part of vector_space_1
   for vector in subspace:
       if vector not in self.matrix:
          return False
   return True
```

We can also check if a given vector space is a subspace by iterating through the original vector space and verifying if each element is in the vector space.

3.1 Identifying The Type of Linear Mapping

There are a handful of different ways to classify a mapping. These can tell us useful properties about a mapping: $\phi: J \to K$

```
Injective if \forall x, y \in J : \phi(x) = \phi(x)
```

```
Injective if \forall x, y \in J : \phi(x) = \phi(y)
Surjective if \phi(J) = K
```

Bijective if both hold

```
Homomorphism if its a linear mapping Isomorphism if \phi: J \to K linear and bijective Endomorphism if \phi: J \to J linear Automorphism if \phi: J \to J linear and bijective Our code to do this is listed below:
```

```
class LinearMapping:
def __init__(self, A, B, mapping=None):
 self.domain_basis = [A[i] for i in
     mp.identify_pivots(mp.transpose(A))]
 self.domain_dim = len(self.domain_basis)
 self.codomain_basis = [B[i] for i in
     mp.identify_pivots(mp.transpose(B))]
 self.codomain_dim = len(self.codomain_basis)
 self.domain = [row[:] for row in A]
 self.codomain = [row[:] for row in B]
 self.mapping = mapping # needs to be a matrix or None
 self.injective = True if mp.rank(self.mapping) ==
     len(self.mapping[0]) else False
 self.surjective = True if mp.rank(self.mapping) ==
     len(self.mapping) else False
 self.bijective = True if (self.injective == True and
     self.surjective == True) else False
 self.homomorphism = True # (by definition, duh)
 self.isomorphism = True if (self.bijective == True) else False
     # homomorphism already satisfied
 self.endomorphism = True if len(A) == len(B) and len(A[0]) ==
     len(B[0]) else False # True if dim(A) == dim(B) and
     len(A[0]) == len(B[0]) else False
 self.automorphism = True if self.endomorphism and
     self.bijective else False
```

3.2 More advanced Basis Change

To find the transformation matrix between basis vectors requires the RREF (Row-Reduced Echelon Form) of the new basis with the older basis vectors, we place the original basis vectors on the left and the desired basis on the right. We then take the right side of the augmented matrix; this can be extended to 2 pairs of bases as shown below:

```
def find_transition(og_base, new_base, columned=False):
, , ,
TAKES:
   original basis of subspace, og_base
   new basis of subspace, new_base
   it is important to pay mind to the shape of these basis
       vectors.
   eg, input [[1,3,4], [1,4,7]] where each sublist is a basis
       vector will be treated as 3 col vectors,
   [1,1], [3,4], and [4,7]. we added the columned flag to help
       with this
, , ,
if columned==False:
   og_base, new_base = transpose(og_base), transpose(new_base)
# we find a transition matrix that changes the coordinates
   expressed from one base to another base
# going from base 1 to base 2
if len(og_base) == len(new_base) and len(og_base[0]) ==
   len(new_base[0]):
   dim_to_take = len(og_base[0]) # we find what the dimensions
       of the transmat will be
   total = append_mat_right(new_base, og_base) # we create an
       augmented matrix with new basis on left and old basis on
       right
   reduced = rref(total) # row reduce, O(n**3)
   transition_matrix = list() # init our blank transition
       matrix
   for row in reduced: # we go through the reduced augmat to
       find what should be added to the transition matrix
       transition_matrix.append(row[-dim_to_take:])
   return transition_matrix
# Full time O(n**3)
```

```
A'_{\phi} = T^{-1}A_{\phi}S

Where \phi: J \to K and ordered bases B = (b_1, ..., b_n)B' = (b'_1, ..., b'_n)

C = (c_1, ..., c_n)C' = (c'_1, ..., c'_n)
```

B is a tuple of the basis vectors in that given space and S represents the mapping of coordinate from B to B'. T represents the mapping of coordinates from C to C' and A_{ϕ} is the Transformation matrix from B to C.

```
def change_transformation_basis(ogtransfmat, ogb1, ogb2,
   tildab1, tildab2, columned=False):
, , ,
TAKES:
   the transformation matrix/linmapping, ogtransfmat, of
       standard form outlined in flowerbox
   the original basis of domain, ogb1
   the original basis of codomain, ogb2
   the new basis of domain, tildab1
   the new basis of codomain, tildab2
   AS WITH find_transition, the base inputs can be finicky,
       the columned flag should help. check that for more
       indepth docs
, , ,
if columned == False: # we make sure the basis inputs are the
   correct form ie they are column vectors
   ogb1, ogb2, tildab1, tildab2 =
       transpose(ogb1),transpose(ogb2),transpose(tildab1),transpose(tildab2)
transition1 = find_transition(tildab1, ogb1) # we find
   transitino matrix from the new domain basis to the old
   domain basis, O(n**3)
transition2 = inverse(find_transition(tildab2, ogb2)) # we find
   inverse of transition matrix from new codomain basis to old
   codomain basis, O(n**3)
return multiply_matrix(multiply_matrix(transition2,
   ogtransfmat), transition1) # we multiply the matrices
   according to formula, O(n**3)
# full time O(n**3)
```

4 More Advanced Matrix Operations

4.1 Solving Homogeneous System

Some matrices contain linearly dependent columns or rows making certain variables "free". We learned about the -1 trick for solving such matrices.

```
def solve_homogeneous(coef_matrix):
# takes matrix of form outlined in flowerbox
# returns a matrix using the minus 1 trick to solve homogeneous
   linear systems
coef_matrix = rref(coef_matrix) # takes the rref of the
   coefficient matrix (system of linear eq) O(n**3)
pivotcols = identify_pivots(coef_matrix) # finds the pivot
   columns of the rref matrix O(n**2)
notpiv_cols = list() # since this function is called because of
   different size dimension matricies, there are going to be
   necessary added piv-columns
for i in range(len(coef_matrix[0])): # iterates through the
   columns O(n)
   if i not in pivotcols: # Iterates through the list of pivot
       notpiv_cols.append(i) # appends the non pivot column to
          the non-pivot col list
if notpiv_cols == []: # if the non-pivot columns are empty (in
   other words had total rank)
   return [0 for i in range(len(coef_matrix[0]))]
if len(coef_matrix) == len(coef_matrix[0]):
   for idx in notpiv_cols:
       coef_matrix[idx][idx] = -1
   coef_matrix = transpose(coef_matrix)
   final = [coef_matrix[idx] for idx in notpiv_cols]
   return final
for idx in notpiv_cols: #Iterate through the non pivot columns
   coef_matrix.insert(idx, [0 if i != idx else -1 for i in
       range(len(coef_matrix[0]))]) # by the minus 1 trick, we
       insert a row with minus 1 in the nth index to preserve
```

```
diagonality
coef_matrix = transpose(coef_matrix) # we transpose the matrix
final = [coef_matrix[idx] for idx in notpiv_cols] # we return
    the solution as a square matrix
return final
# O(n**3)
```

5 Projections

- 5.1 Solving Homogeneous Systems
- 5.2 Finding Eigenvalues and Eigenvectors Of A Matrix
- 5.3 Finding the Eiegendecomposition
- 5.4 Finding the Singular Value Decomposition