AntiMatter

Adam Levin-Delson

Copyright © Copyright©1995 Adam Levin-Delson. All Rights Reserved

COLLABORATORS							
	TITLE :						
AntiMatter							
ACTION	NAME	DATE	SIGNATURE				
WRITTEN BY	Adam Levin-Delson	March 15, 2022					

REVISION HISTORY							
NUMBER	DATE	DESCRIPTION	NAME				

Contents

1	Anti	Matter	1
	1.1	AntiMatter User Manual	1
	1.2	features	1
	1.3	instructions	2
	1.4	starting antimatter	2
	1.5	startup wb	2
	1.6	startup shell	3
	1.7	startup priority	3
	1.8	firing rays	3
	1.9	ray travel	3
	1.10	unimpeded flight	4
	1.11	absorption	4
	1.12	deflection	4
	1.13	reflection	4
	1.14	marking choices	4
	1.15	scoring	5
	1.16	score calculation	5
	1.17	menus	5
	1.18	project menu	5
	1.19	prefs menu	6
	1.20	guess menu	6
	1.21	tips	6
	1.22	tips preferences	6
	1.23	tips beginner games	6
	1.24	tips deflections	6
	1.25	tips impossibilities	7
	1.26	help	7
	1.27	bug reports	7
	1.28	source code	7
	1.29	contacting the author	7
	1.30	acknowledgements	8
	1.31	release history	8

Chapter 1

AntiMatter

1.1 AntiMatter User Manual

AntiMatter

Copyright © 1988-1995 Adam Levin-Delson All Rights Reserved

Permission is hereby granted to distribute the AntiMatter executable and documentation via electronic means and in Amiga software collections for which no charge or only a small amount of money more than production costs is charged.

AntiMatter is a single-player strategy game in which the computer hides a number of atoms in a chamber. You must determine the locations of the atoms by firing plasma rays into the chamber and observing where the rays (which are influenced by the atoms) emerge.

Features - Features of this implementation.

Instructions - How to play AntiMatter.

Bug reports - How to report bugs to the author.

Source code - How to obtain the Source code.

Acknowledgements - Acknowledgements and credits.

Release history - Describes the various versions of AntiMatter.

Documentation is available during gameplay at the press of the Help key. To enable this feature, assign "AM:" to the directory where AntiMatter.doc can be found.

1.2 features

AntiMatter Features

Stereo sound

- Four-channel stereo sound follows the action!

- Conforms to the Amiga's "channel-locking" convention, allowing channels to be stolen by higher priority requests, and later returned to the original owner.

Animated Pointer

- 30 fps animated pointer using a proprietary vertical-blank interrupt handler.

- Multiple instantiations of AntiMatter share the animating routine.

Configurable

- The player can choose how large a chamber to use (6x6 through 14x14) as well as how many atoms will be hidden (3 through 8). (See Preferences .)

OS-sensitive

- Runs under AmigaOS 1.3, 2.x and 3.x and on all 680x0 processors.

Low memory usage

- Opens a low-res screen and uses a "shading" pattern to extend the color palette.

- Chip RAM audio samples are shared for the various sound effects.

Well tested

- Developed using Enforcer and Mungwall.

1.3 instructions

AntiMatter Instructions

A new game is set up when Antimatter is first invoked. Changing the configuration of the board (see Preferences) will also set up a new game, as will selecting "New game" from the "Project" menu.

Starting Antimatter Firing rays Ray travel Marking choices Scoring Menus Online Help Tips

1.4 starting antimatter

Instructions - Starting Antimatter

Antimatter can be started from either the Workbench (by double-clicking on the Antimatter icon) or from the shell. Game preferences can be specified from either interface.

Workbench startup

Shell startup

1.5 startup wb

Starting AntiMatter - Workbench

The number of atoms and size of the chamber can be set in AntiMatter's icon (using "Information" from the Workbench) with "ATOMS=" and "SIZE=" tooltypes. An additional ToolType parameter, "SOUNDPRI=" can be used to set the priority of Anti-Matter's sound effects. If SOUNDPRI is set to the special value "999", AntiMatter will not generate any sounds.

Valid numbers of atoms range from 3 to 8; chamber size from 6 (6x6) to 14 (14x14).

The default values for the tooltypes are as follows:

ATOMS=4

SIZE=8

SOUNDPRI=127

1.6 startup shell

Starting AntiMatter - Shell

The number of atoms and size of the chamber can be specified when starting AntiMatter from the Command Line (Shell). An additional argument can be used to set the priority of AntiMatter's sound effects. If SOUNDPRI is set to the special value "999", AntiMatter will not generate any sounds.

Valid numbers of atoms range from 3 to 8; chamber size from 6 (6x6) to 14 (14x14).

The syntax of the command depends upon the version of the operating system installed on your Amiga.

Newer versions of AmigaDOS (specifically, version 2.04 and later) use the "ReadArgs" syntax:

Antimatter ATOMS=4 SIZE=8 SOUNDPRI=127

Older versions use a Unix-style syntax:

AntiMatter -a4 -s8 -p127

Both formats shown above illustrate the default AntiMatter values.

1.7 startup priority

Starting AntiMatter - Sound Priority

AntiMatter allocates all four Amiga audio channels to create action-specific stereo sound. A priority level of 127 will prevent any other application from stealing AntiMatter's audio channels. Anything less than 127 (down to -128) will allow applications with a higher audio priority to steal the channels. AntiMatter will attempt to reallocate stolen channels as soon as they are freed.

1.8 firing rays

Instructions - Firing Rays

Fire a ray by positioning the laser (cursor) at one of the chamber entrances (the squares along the chamber edges) and clicking the left mouse button.

The entrance (and exit, if the ray ultimately exited the chamber) will then be indicated with a colored marker (described in Ray travel) determined by how the chamber's atoms influenced the ray.

1.9 ray travel

Instructions - Ray travel

A ray will normally travel in a straight line (an Unimpeded flight, the ray passes straight through the chamber) unless acted upon by one or more atoms.

There are two primary ways in which a ray can be effected by atoms:

Absorption - The ray is absorbed by an atom in the chamber.

Deflection - The ray is caused to veer 90° from a straight line.

There are two special cases of Deflections known as Reflections .

Note that the influence of multiple atoms in the chamber can combine to yield complex results.

1.10 unimpeded flight

Ray travel - Unimpeded flight

If there are no atoms directly in the ray's path, nor along the left or right of the path, the ray will pass straight through the chamber, exiting on the opposite side.

The entrance and exit locations will be indicated with like-color markers.

1.11 absorption

Ray travel - Absorption

If a ray is fired directly into an atom, it is absorbed, and does not exit the chamber.

The square where the ray entered the chamber will be indicated with a black marker.

1.12 deflection

Ray travel - Deflection

If a ray would pass immediately alongside an atom the ray will be deflected away from the atom at a right angle to its original path.

If the ray would have passed the atom on the atom's right side, the ray is deflected to the right.

If the ray would have passed on the atom's left side, it is deflected to the left.

Note that the deflection occurs one square *before* the ray and the atom would be next to each other. The entrance and exit locations will be indicated with like-color markers.

There are two important conditions that result in a special type of deflection known as a Reflection .

1.13 reflection

Ray travel - Reflection

If the deflecting atom is alongside the square at which the ray would enter the chamber, the ray is unable to be deflected, and is instead reflected directly back out.

This reflection is indicated with a white marker at the attempted entrance point.

A reflection also occurs if there are two atoms straddling the ray's path (ie they are exactly one square apart and the ray is traveling down the path between them). The ray is deflected both right and left at the same time, but cannot overcome either atom, and is instead reflected back along the path from which it came.

The ray emerges from the same location it entered, and the reflection is indicated with a white marker.

1.14 marking choices

Instructions - Marking choices

As you fire rays into the chamber, and observe the results, you will start to form an idea of where the hidden atoms are.

You can mark your suspected locations by positioning the cursor over the desired location in the center grid of the chamber and clicking the left mouse button, creating a white marker.

Clicking on the same location again will remove the marker. While you may place as many markers in the center grid of the chamber as you like, you will not be able to score your guesses if you have more markers than there are atoms hidden in the chamber.

1.15 scoring

Instructions - Scoring

During game play, you may place as many markers in the center grid of the chamber as you like. In order to calculate your score, however, you may have no more markers than there are atoms hidden in the chamber.

Once you believe that you have correctly guessed the location of all the atoms in the chamber, (as indicated by your white markers in the center grid), choose the menu item "Guess/How did I do?".

If the "How did I do?" menu choice is ghosted (disabled), you have too many markers on the center grid of the chamber. Simply click on the unneeded markers until you have as many (or fewer) markers as there are atoms hidden in the chamber.

To calculate your score, the actual locations of the atoms will be displayed. Atoms which you correctly guessed will be marked with a checkerboard marker, atoms which you missed will be solid red, and atoms which you thought existed but didn't are left in white.

Your score is calculated from the number of rays fired, and the number of atoms correctly guessed. The lower your score the better.

1.16 score calculation

Instructions - Score calculation

The formula used for calculating the score is:

(5*(A-C)) + R

where

A is the number of atoms embedded in the chamber.

C is the number of atoms correctly identified

with markers in the center of the chamber.

R is the number of ray markers along the edges

of the chamber.

1.17 menus

Instructions - Menus AntiMatter has three main menus: Project menu Prefs menu Guess menu

1.18 project menu

Menus - Project Menu

The "Project/New Game" menu item (Keyboard shortcut "Right-Amiga N") will begin a new game with the selected game parameters of chamber size and number of atoms.

The "Project/About..." menu item ("Right-Amiga A") gives credit for the original board game upon which AntiMatter is based.

The "Project/Quit" menu item ("Right-Amiga Q") will exit the game.

1.19 prefs menu

Menus - Prefs Menu

There are several menu selections that will help you tailor AntiMatter to your desired level of difficulty. The "Prefs/Atoms" and "Prefs/Size of Chamber" menu items will allow you to set the number of atoms and the size of the chamber. The currently selected values are displayed in a "ghosted" or dimmed manner. As soon as you change one of these parameters, a new game will be created using the new value(s). You can choose from both the "Atoms" and "Size of Box" menus without releasing the "menu" (right) mouse button in between. This is known as extended-selecting the items.

These preferences can also be specified when AntiMatter is invoked from either the Workbench or Shell .

1.20 guess menu

Menus - Guess Menu

The "Guess/Clear Markers" and "Guess/Set Markers" menu items (Keyboard shortcuts "Right-Amiga C" and "Right-Amiga S", respectively) will allow you to clear or set all the center grid markers at once.

The "Guess/How did I do?" menu item ("Right-Amiga T") will tally up your score. You will be unable to select this item if there are too many "guess" markers on the center grid.

1.21 tips

Instructions - Tips Preferences Beginner games Deflections Impossibilities

1.22 tips preferences

Tips - Preferences

The default size of 8x8 and 4 atoms is a good starting point, but feel free to make the chamber smaller or the atoms fewer when learning how the rays are effected by the atoms.

1.23 tips beginner games

Tips - Beginner Games

To help in understanding how rays are effected by the atoms, you may choose to fire rays into all possible entrances, then select the "Guess/How did I Do" menu to see the locations of the atoms, and then study the path of all the rays.

1.24 tips deflections

Tips - Deflections

Keep in mind that Deflections occur one square before the ray would be alongside an atom.

1.25 tips impossibilities

Tips - Impossibilities

There are configurations of atoms ("molecules") which cannot be completely determined even if rays are fired into all chamber entrances.

The likelihood of such molecules increases with the number of atoms embedded in the chamber.

1.26 help

Instructions - Online Help

Pressing the "Help" key during game play will interrupt the action and call up "AmigaGuide" to display this documentation.

AmigaGuide will be instructed to look in the same directory that AntiMatter is located in for "AntiMatter.guide".

Should either AmigaGuide or the guide file not be found, AntiMatter will attempt to call up "More" (actually Sys:Utilities/More) on "AM:AntiMatter.doc".

1.27 bug reports

Bug reports

Bug Reports are encouraged, and may be sent electronically or by mail to the author .

The first person to report each reproducible bug will receive a corrected copy as soon as it is available.

1.28 source code

Source code

SAS 'C' (v6.51) source code for AntiMatter is available from the author for US\$10 (this includes postage in the United States; peoples of all other countries [unite!] must include appropriate postage to pay for the shipment of one diskette from the US to your country).

680x0 assembler source code for AnimPtr, the sharable vertical-interrupt handler for pointer animation is available for US\$5.

Note that the source code for AntiMatter and AnimPtr are not redistributable. They may, however, be used as the basis for other programs, which may be distributed as you see fit, so long as an acknowledgement of my original source is clearly given.

1.29 contacting the author

AntiMatter - Contacting the author

Contact me to report bugs found in AntiMatter, or to obtain a copy of the source code.

My mail address is:

Adam Levin-Delson

22325 53rd Ave SE

Bothell, WA 98021 USA

My E-Mail address is: aklevin@bix.com

1.30 acknowledgements

AntiMatter - Acknowledgements

AntiMatter is based upon the board game "Black Box", by Eric Solomon. It was originally produced by Waddingtons House of Games in the United Kingdom, and then by Parker Brothers in the United States of America.

AntiMatter is Copyright © 1988-1995 Adam Levin-Delson All Rights Reserved

Many thanks to the following people:

Bernie Cosell and Bruce Barnett, for their kind help in providing information and materials pertaining to the two board-game versions of "Black Box".

Michael Sinz, Peter Cherna, Spencer Shanson and Carolyn Scheppner, all of (at the time) Commodore Business Machines.

Peter Kunath for audio channel-sharing example code.

Boyd & Gian Edmondson for beta-testing.

Toby Douglass for bug-reporting.

1.31 release history

AntiMatter - Release history

AntiMatter's version number can be found in the information provided by the "About..." menu selection as well as from the AmigaDOS "Version" command.

Version Date Comments

1.0 1988 Working (audioless) version. Not distributed.

1.XX 1993 - 1994 Took advantage of AmigaOS 2.x/3.x features.

Added animated pointer.

Added sound routines.

2.0 06 Jan 1995 First full release.

2.1 25 Jan 1995 Fixed bug preventing Euro36 screens from opening. (Toby Douglass found this.)